

Alternatively, techniques described for the production of single chain antibodies (U.S. Patent No. 4,946,778; Bird, Science 242:423- 42 (1988); Huston et al., Proc. Natl. Acad. Sci. USA 85:5879-5883 (1988); and Ward et al., Nature 334:544-54 (1989)) can be adapted to produce single chain antibodies. Single chain antibodies are formed by linking the heavy and light chain fragments of the Fv region via an amino acid bridge, resulting in a single chain polypeptide. Techniques for the assembly of functional Fv fragments in E. coli may also be used (Skerra et al., Science 242:1038- 1041 (1988)).

Methods of Producing Antibodies

The antibodies of the invention can be produced by any method known in the art for the synthesis of antibodies, in particular, by chemical synthesis or preferably, by recombinant expression techniques.

Recombinant expression of an antibody of the invention, or fragment, derivative or analog thereof, (e.g., a heavy or light chain of an antibody of the invention or a single chain antibody of the invention), requires construction of an expression vector containing a polynucleotide that encodes the antibody. Once a polynucleotide encoding an antibody molecule or a heavy or light chain of an antibody, or portion thereof (preferably containing the heavy or light chain variable domain), of the invention has been obtained, the vector for the production of the antibody molecule may be produced by recombinant DNA technology using techniques well known in the art. Thus, methods for preparing a protein by expressing a polynucleotide containing an antibody encoding nucleotide sequence are described herein. Methods which are well known to those skilled in the art can be used to construct expression vectors containing antibody coding sequences and appropriate transcriptional and translational control signals. These methods include, for example, in vitro recombinant DNA techniques, synthetic techniques, and in vivo genetic recombination. The invention, thus, provides replicable vectors comprising a nucleotide sequence encoding an antibody molecule of the invention, or a heavy or light chain thereof, or a heavy or light chain variable domain, operably linked to a promoter. Such vectors may include the nucleotide sequence encoding the constant region of the antibody molecule (see, e.g., PCT Publication WO 86/05807; PCT Publication WO 89/01036; and U.S. Patent No. 5,122,464) and the variable domain of the antibody may be cloned into such a vector for expression of the entire heavy or light chain.

The expression vector is transferred to a host cell by conventional techniques and the transfected cells are then cultured by conventional techniques to produce an antibody of the invention. Thus, the invention includes host cells containing a polynucleotide encoding an antibody of the invention, or a heavy or light chain thereof, or a single chain antibody of the invention, operably linked to a heterologous promoter. In preferred embodiments for the expression of double-chained antibodies, vectors encoding both the heavy and light chains may be co-expressed in the host cell for expression of the entire immunoglobulin molecule, as detailed below.

A variety of host-expression vector systems may be utilized to express the antibody molecules of the invention. Such host-expression systems represent vehicles by which the coding sequences of interest may be produced and subsequently purified, but also represent cells which may, when transformed or transfected with the appropriate nucleotide coding sequences, express an antibody molecule of the invention in situ. These include but are not limited to microorganisms such as bacteria (e.g., *E. coli*, *B. subtilis*) transformed with recombinant bacteriophage DNA, plasmid DNA or cosmid DNA expression vectors containing antibody coding sequences; yeast (e.g., *Saccharomyces*, *Pichia*) transformed with recombinant yeast expression vectors containing antibody coding sequences; insect cell systems infected with recombinant virus expression vectors (e.g., baculovirus) containing antibody coding sequences; plant cell systems infected with recombinant virus expression vectors (e.g., cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or transformed with recombinant plasmid expression vectors (e.g., Ti plasmid) containing antibody coding sequences; or mammalian cell systems (e.g., COS, CHO, BHK, 293, 3T3 cells) harboring recombinant expression constructs containing promoters derived from the genome of mammalian cells (e.g., metallothionein promoter) or from mammalian viruses (e.g., the adenovirus late promoter; the vaccinia virus 7.5K promoter). Preferably, bacterial cells such as *Escherichia coli*, and more preferably, eukaryotic cells, especially for the expression of whole recombinant antibody molecule, are used for the expression of a recombinant antibody molecule. For example, mammalian cells such as Chinese hamster ovary cells (CHO), in conjunction with a vector such as the major intermediate early gene promoter element from human cytomegalovirus is an effective expression system for antibodies (Foecking et al., Gene 45:101 (1986); Cockett et al., Bio/Technology 8:2 (1990)).

In bacterial systems, a number of expression vectors may be advantageously selected depending upon the use intended for the antibody molecule being expressed. For example, when a large quantity of such a protein is to be produced, for the generation of pharmaceutical compositions of an antibody molecule, vectors which direct the expression of high levels of fusion protein products that are readily purified may be desirable. Such vectors include, but are not limited, to the *E. coli* expression vector pUR278 (Ruther et al., EMBO J. 2:1791 (1983)), in which the antibody coding sequence may be ligated individually into the vector in frame with the lac Z coding region so that a fusion protein is produced; pIN vectors (Inouye & Inouye, Nucleic Acids Res. 13:3101-3109 (1985); Van Heeke & Schuster, J. Biol. Chem. 24:5503-5509 (1989)); and the like. pGEX vectors may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such fusion proteins are soluble and can easily be purified from lysed cells by adsorption and binding to matrix glutathione-agarose beads followed by elution in the presence of free glutathione. The pGEX vectors are designed to include thrombin or factor Xa protease cleavage sites so that the cloned target gene product can be released from the GST moiety.

In an insect system, Autographa californica nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes. The virus grows in *Spodoptera frugiperda* cells. The antibody coding sequence may be cloned individually into non-essential regions (for example the polyhedrin gene) of the virus and placed under control of an AcNPV promoter (for example the polyhedrin promoter).

In mammalian host cells, a number of viral-based expression systems may be utilized. In cases where an adenovirus is used as an expression vector, the antibody coding sequence of interest may be ligated to an adenovirus transcription/translation control complex, e.g., the late promoter and tripartite leader sequence. This chimeric gene may then be inserted in the adenovirus genome by in vitro or in vivo recombination. Insertion in a non-essential region of the viral genome (e.g., region E1 or E3) will result in a recombinant virus that is viable and capable of expressing the antibody molecule in infected hosts. (e.g., see Logan & Shenk, Proc. Natl. Acad. Sci. USA 81:355-359 (1984)). Specific initiation signals may also be required for efficient translation of inserted antibody coding sequences. These signals include the ATG initiation codon and adjacent sequences. Furthermore, the initiation codon must be in phase with the reading frame of the desired coding sequence to ensure translation of the entire insert. These exogenous translational control signals and initiation codons can

be of a variety of origins, both natural and synthetic. The efficiency of expression may be enhanced by the inclusion of appropriate transcription enhancer elements, transcription terminators, etc. (see Bittner et al., *Methods in Enzymol.* 153:51-544 (1987)).

In addition, a host cell strain may be chosen which modulates the expression of the
5 inserted sequences, or modifies and processes the gene product in the specific fashion desired. Such modifications (e.g., glycosylation) and processing (e.g., cleavage) of protein products may be important for the function of the protein. Different host cells have characteristic and specific mechanisms for the post-translational processing and modification of proteins and gene products. Appropriate cell lines or host systems can be chosen to
10 ensure the correct modification and processing of the foreign protein expressed. To this end, eukaryotic host cells which possess the cellular machinery for proper processing of the primary transcript, glycosylation, and phosphorylation of the gene product may be used. Such mammalian host cells include but are not limited to CHO, VERY, BHK, HeLa, COS, MDCK, 293, 3T3, WI38, and in particular, breast cancer cell lines such as, for example,
15 BT483, Hs578T, HTB2, BT20 and T47D, and normal mammary gland cell line such as, for example, CRL7030 and Hs578Bst.

For long-term, high-yield production of recombinant proteins, stable expression is preferred. For example, cell lines which stably express the antibody molecule may be engineered. Rather than using expression vectors which contain viral origins of replication,
20 host cells can be transformed with DNA controlled by appropriate expression control elements (e.g., promoter, enhancer, sequences, transcription terminators, polyadenylation sites, etc.), and a selectable marker. Following the introduction of the foreign DNA, engineered cells may be allowed to grow for 1-2 days in an enriched media, and then are switched to a selective media. The selectable marker in the recombinant plasmid confers
25 resistance to the selection and allows cells to stably integrate the plasmid into their chromosomes and grow to form foci which in turn can be cloned and expanded into cell lines. This method may advantageously be used to engineer cell lines which express the antibody molecule. Such engineered cell lines may be particularly useful in screening and evaluation of compounds that interact directly or indirectly with the antibody molecule.

30 A number of selection systems may be used, including but not limited to the herpes simplex virus thymidine kinase (Wigler et al., *Cell* 11:223 (1977)), hypoxanthine-guanine phosphoribosyltransferase (Szybalska & Szybalski, *Proc. Natl. Acad. Sci. USA* 48:202

(1992)), and adenine phosphoribosyltransferase (Lowy et al., Cell 22:817 (1980)) genes can be employed in tk-, hgp^rt- or ap^rt- cells, respectively. Also, antimetabolite resistance can be used as the basis of selection for the following genes: dhfr, which confers resistance to methotrexate (Wigler et al., Natl. Acad. Sci. USA 77:357 (1980); O'Hare et al., Proc. Natl. Acad. Sci. USA 78:1527 (1981)); gpt, which confers resistance to mycophenolic acid (Mulligan & Berg, Proc. Natl. Acad. Sci. USA 78:2072 (1981)); neo, which confers resistance to the aminoglycoside G-418 Clinical Pharmacy 12:488-505; Wu and Wu, Biotherapy 3:87-95 (1991); Tolstoshev, Ann. Rev. Pharmacol. Toxicol. 32:573-596 (1993); Mulligan, Science 260:926-932 (1993); and Morgan and Anderson, Ann. Rev. Biochem. 62:191-217 (1993); May, 1993, TIB TECH 11(5):155-215); and hyg^r, which confers resistance to hygromycin (Santerre et al., Gene 30:147 (1984)). Methods commonly known in the art of recombinant DNA technology may be routinely applied to select the desired recombinant clone, and such methods are described, for example, in Ausubel et al. (eds.), Current Protocols in Molecular Biology, John Wiley & Sons, NY (1993); Kriegler, Gene Transfer and Expression, A Laboratory Manual, Stockton Press, NY (1990); and in Chapters 12 and 13, Dracopoli et al. (eds), Current Protocols in Human Genetics, John Wiley & Sons, NY (1994); Colberre-Garapin et al., J. Mol. Biol. 150:1 (1981), which are incorporated by reference herein in their entireties.

The expression levels of an antibody molecule can be increased by vector amplification (for a review, see Bebbington and Hentschel, The use of vectors based on gene amplification for the expression of cloned genes in mammalian cells in DNA cloning, Vol.3. (Academic Press, New York, 1987)). When a marker in the vector system expressing antibody is amplifiable, increase in the level of inhibitor present in culture of host cell will increase the number of copies of the marker gene. Since the amplified region is associated with the antibody gene, production of the antibody will also increase (Crouse et al., Mol. Cell. Biol. 3:257 (1983)).

The host cell may be co-transfected with two expression vectors of the invention, the first vector encoding a heavy chain derived polypeptide and the second vector encoding a light chain derived polypeptide. The two vectors may contain identical selectable markers which enable equal expression of heavy and light chain polypeptides. Alternatively, a single vector may be used which encodes, and is capable of expressing, both heavy and light chain polypeptides. In such situations, the light chain should be placed before the heavy chain to

avoid an excess of toxic free heavy chain (Proudfoot, Nature 322:52 (1986); Kohler, Proc. Natl. Acad. Sci. USA 77:2197 (1980)). The coding sequences for the heavy and light chains may comprise cDNA or genomic DNA.

Once an antibody molecule of the invention has been produced by an animal, 5 chemically synthesized, or recombinantly expressed, it may be purified by any method known in the art for purification of an immunoglobulin molecule, for example, by chromatography (e.g., ion exchange, affinity, particularly by affinity for the specific antigen after Protein A, and sizing column chromatography), centrifugation, differential solubility, or by any other standard technique for the purification of proteins. In addition, the antibodies of 10 the present invention or fragments thereof can be fused to heterologous polypeptide sequences described herein or otherwise known in the art, to facilitate purification.

The present invention encompasses antibodies recombinantly fused or chemically conjugated (including both covalently and non-covalently conjugations) to a polypeptide (or 15 portion thereof, preferably at least 10, 20, 30, 40, 50, 60, 70, 80, 90 or 100 amino acids of the polypeptide) of the present invention to generate fusion proteins. The fusion does not necessarily need to be direct, but may occur through linker sequences. The antibodies may be specific for antigens other than polypeptides (or portion thereof, preferably at least 10, 20, 30, 40, 50, 60, 70, 80, 90 or 100 amino acids of the polypeptide) of the present invention. For 20 example, antibodies may be used to target the polypeptides of the present invention to particular cell types, either in vitro or in vivo, by fusing or conjugating the polypeptides of the present invention to antibodies specific for particular cell surface receptors. Antibodies fused or conjugated to the polypeptides of the present invention may also be used in in vitro immunoassays and purification methods using methods known in the art. See e.g., Harbor et al., supra, and PCT publication WO 93/21232; EP 439,095; Naramura et al., Immunol. Lett. 25 39:91-99 (1994); U.S. Patent 5,474,981; Gillies et al., PNAS 89:1428-1432 (1992); Fell et al., J. Immunol. 146:2446-2452(1991), which are incorporated by reference in their entireties.

The present invention further includes compositions comprising the polypeptides of the present invention fused or conjugated to antibody domains other than the variable regions. For example, the polypeptides of the present invention may be fused or conjugated to an 30 antibody Fc region, or portion thereof. The antibody portion fused to a polypeptide of the present invention may comprise the constant region, hinge region, CH1 domain, CH2 domain, and CH3 domain or any combination of whole domains or portions thereof. The

polypeptides may also be fused or conjugated to the above antibody portions to form multimers. For example, Fc portions fused to the polypeptides of the present invention can form dimers through disulfide bonding between the Fc portions. Higher multimeric forms can be made by fusing the polypeptides to portions of IgA and IgM. Methods for fusing or
5 conjugating the polypeptides of the present invention to antibody portions are known in the art. See, e.g., U.S. Patent Nos. 5,336,603; 5,622,929; 5,359,046; 5,349,053; 5,447,851; 5,112,946; EP 307,434; EP 367,166; PCT publications WO 96/04388; WO 91/06570; Ashkenazi et al., Proc. Natl. Acad. Sci. USA 88:10535-10539 (1991); Zheng et al., J. Immunol. 154:5590-5600 (1995); and Vil et al., Proc. Natl. Acad. Sci. USA 89:11337-
10 11341(1992) (said references incorporated by reference in their entireties).

As discussed, supra, the polypeptides corresponding to a polypeptide, polypeptide fragment, or a variant of SEQ ID NO:Y may be fused or conjugated to the above antibody portions to increase the in vivo half life of the polypeptides or for use in immunoassays using methods known in the art. Further, the polypeptides corresponding to SEQ ID NO:Y may be
15 fused or conjugated to the above antibody portions to facilitate purification. One reported example describes chimeric proteins consisting of the first two domains of the human CD4-polypeptide and various domains of the constant regions of the heavy or light chains of mammalian immunoglobulins. (EP 394,827; Traunecker et al., Nature 331:84-86 (1988). The polypeptides of the present invention fused or conjugated to an antibody having
20 disulfide-linked dimeric structures (due to the IgG) may also be more efficient in binding and neutralizing other molecules, than the monomeric secreted protein or protein fragment alone. (Fountoulakis et al., J. Biochem. 270:3958-3964 (1995)). In many cases, the Fc part in a fusion protein is beneficial in therapy and diagnosis, and thus can result in, for example, improved pharmacokinetic properties. (EP A 232,262). Alternatively, deleting the Fc part
25 after the fusion protein has been expressed, detected, and purified, would be desired. For example, the Fc portion may hinder therapy and diagnosis if the fusion protein is used as an antigen for immunizations. In drug discovery, for example, human proteins, such as hIL-5, have been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists of hIL-5. (See, Bennett et al., J. Molecular Recognition 8:52-58 (1995);
30 Johanson et al., J. Biol. Chem. 270:9459-9471 (1995).

Moreover, the antibodies or fragments thereof of the present invention can be fused to marker sequences, such as a peptide to facilitate purification. In preferred embodiments, the

marker amino acid sequence is a hexa-histidine peptide, such as the tag provided in a pQE vector (QIAGEN, Inc., 9259 Eton Avenue, Chatsworth, CA, 91311), among others, many of which are commercially available. As described in Gentz et al., Proc. Natl. Acad. Sci. USA 86:821-824 (1989), for instance, hexa-histidine provides for convenient purification of the fusion protein. Other peptide tags useful for purification include, but are not limited to, the "HA" tag, which corresponds to an epitope derived from the influenza hemagglutinin protein (Wilson et al., Cell 37:767 (1984)) and the "flag" tag.

The present invention further encompasses antibodies or fragments thereof conjugated to a diagnostic or therapeutic agent. The antibodies can be used diagnostically to, for example, monitor the development or progression of a tumor as part of a clinical testing procedure to, e.g., determine the efficacy of a given treatment regimen. Detection can be facilitated by coupling the antibody to a detectable substance. Examples of detectable substances include various enzymes, prosthetic groups, fluorescent materials, luminescent materials, bioluminescent materials, radioactive materials, positron emitting metals using various positron emission tomographies, and nonradioactive paramagnetic metal ions. The detectable substance may be coupled or conjugated either directly to the antibody (or fragment thereof) or indirectly, through an intermediate (such as, for example, a linker known in the art) using techniques known in the art. See, for example, U.S. Patent No. 4,741,900 for metal ions which can be conjugated to antibodies for use as diagnostics according to the present invention. Examples of suitable enzymes include horseradish peroxidase, alkaline phosphatase, beta-galactosidase, or acetylcholinesterase; examples of suitable prosthetic group complexes include streptavidin/biotin and avidin/biotin; examples of suitable fluorescent materials include umbelliferone, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine fluorescein, dansyl chloride or phycoerythrin; an example of a luminescent material includes luminol; examples of bioluminescent materials include luciferase, luciferin, and aequorin; and examples of suitable radioactive material include ^{125}I , ^{131}I , ^{111}In or ^{99}Tc .

Further, an antibody or fragment thereof may be conjugated to a therapeutic moiety such as a cytotoxin, e.g., a cytostatic or cytocidal agent, a therapeutic agent or a radioactive metal ion, e.g., alpha-emitters such as, for example, ^{213}Bi . A cytotoxin or cytotoxic agent includes any agent that is detrimental to cells. Examples include paclitaxol, cytochalasin B, gramicidin D, ethidium bromide, emetine, mitomycin, etoposide, tenoposide, vincristine,

vinblastine, colchicin, doxorubicin, daunorubicin, dihydroxy anthracin dione, mitoxantrone, mithramycin, actinomycin D, 1-dehydrotestosterone, glucocorticoids, procaine, tetracaine, lidocaine, propranolol, and puromycin and analogs or homologs thereof. Therapeutic agents include, but are not limited to, antimetabolites (e.g., methotrexate, 6-mercaptopurine, 6-thioguanine, cytarabine, 5-fluorouracil decarbazine), alkylating agents (e.g., mechlorethamine, thioepa chlorambucil, melphalan, carmustine (BSNU) and lomustine (CCNU), cyclophosphamide, busulfan, dibromomannitol, streptozotocin, mitomycin C, and cis- dichlorodiamine platinum (II) (DDP) cisplatin), anthracyclines (e.g., daunorubicin (formerly daunomycin) and doxorubicin), antibiotics (e.g., dactinomycin (formerly actinomycin), bleomycin, mithramycin, and anthramycin (AMC)), and anti-mitotic agents (e.g., vincristine and vinblastine).

The conjugates of the invention can be used for modifying a given biological response, the therapeutic agent or drug moiety is not to be construed as limited to classical chemical therapeutic agents. For example, the drug moiety may be a protein or polypeptide possessing a desired biological activity. Such proteins may include, for example, a toxin such as abrin, ricin A, pseudomonas exotoxin, or diphtheria toxin; a protein such as tumor necrosis factor, α -interferon, β -interferon, nerve growth factor, platelet derived growth factor, tissue plasminogen activator, an apoptotic agent, e.g., TNF- α , TNF- β , AIM I (See, International Publication No. WO 97/33899), AIM II (See, International Publication No. WO 97/34911), Fas Ligand (Takahashi *et al.*, *Int. Immunol.*, 6:1567-1574 (1994)), VEGI (See, International Publication No. WO 99/23105), a thrombotic agent or an anti- angiogenic agent, e.g., angiostatin or endostatin; or, biological response modifiers such as, for example, lymphokines, interleukin-1 ("IL-1"), interleukin-2 ("IL-2"), interleukin-6 ("IL-6"), granulocyte macrophage colony stimulating factor ("GM-CSF"), granulocyte colony stimulating factor ("G-CSF"), or other growth factors.

Antibodies may also be attached to solid supports, which are particularly useful for immunoassays or purification of the target antigen. Such solid supports include, but are not limited to, glass, cellulose, polyacrylamide, nylon, polystyrene, polyvinyl chloride or polypropylene.

Techniques for conjugating such therapeutic moiety to antibodies are well known, see, e.g., Arnon *et al.*, "Monoclonal Antibodies For Immunotargeting Of Drugs In Cancer Therapy", in *Monoclonal Antibodies And Cancer Therapy*, Reisfeld *et al.* (eds.), pp. 243-56

(Alan R. Liss, Inc. 1985); Hellstrom et al., "Antibodies For Drug Delivery", in Controlled Drug Delivery (2nd Ed.), Robinson et al. (eds.), pp. 623-53 (Marcel Dekker, Inc. 1987); Thorpe, "Antibody Carriers Of Cytotoxic Agents In Cancer Therapy: A Review", in Monoclonal Antibodies '84: Biological And Clinical Applications, Pinchera et al. (eds.), pp. 475-506 (1985); "Analysis, Results, And Future Prospective Of The Therapeutic Use Of Radiolabeled Antibody In Cancer Therapy", in Monoclonal Antibodies For Cancer Detection And Therapy, Baldwin et al. (eds.), pp. 303-16 (Academic Press 1985), and Thorpe et al., "The Preparation And Cytotoxic Properties Of Antibody-Toxin Conjugates", Immunol. Rev. 62:119-58 (1982).

Alternatively, an antibody can be conjugated to a second antibody to form an antibody heteroconjugate as described by Segal in U.S. Patent No. 4,676,980, which is incorporated herein by reference in its entirety.

An antibody, with or without a therapeutic moiety conjugated to it, administered alone or in combination with cytotoxic factor(s) and/or cytokine(s) can be used as a therapeutic.

Immunophenotyping

The antibodies of the invention may be utilized for immunophenotyping of cell lines and biological samples. The translation product of the gene of the present invention may be useful as a cell specific marker, or more specifically as a cellular marker that is differentially expressed at various stages of differentiation and/or maturation of particular cell types. Monoclonal antibodies directed against a specific epitope, or combination of epitopes, will allow for the screening of cellular populations expressing the marker. Various techniques can be utilized using monoclonal antibodies to screen for cellular populations expressing the marker(s), and include magnetic separation using antibody-coated magnetic beads, "panning" with antibody attached to a solid matrix (i.e., plate), and flow cytometry (See, e.g., U.S. Patent 5,985,660; and Morrison *et al.*, *Cell*, 96:737-49 (1999)).

These techniques allow for the screening of particular populations of cells, such as might be found with hematological malignancies (i.e. minimal residual disease (MRD) in acute leukemic patients) and "non-self" cells in transplantations to prevent Graft-versus-Host Disease (GVHD). Alternatively, these techniques allow for the screening of hematopoietic stem and progenitor cells capable of undergoing proliferation and/or differentiation, as might be found in human umbilical cord blood.

Assays For Antibody Binding

The antibodies of the invention may be assayed for immunospecific binding by any method known in the art. The immunoassays which can be used include but are not limited to competitive and non-competitive assay systems using techniques such as western blots, radioimmunoassays, ELISA (enzyme linked immunosorbent assay), "sandwich" immunoassays, immunoprecipitation assays, precipitin reactions, gel diffusion precipitin reactions, immunodiffusion assays, agglutination assays, complement-fixation assays, immunoradiometric assays, fluorescent immunoassays, protein A immunoassays, to name but a few. Such assays are routine and well known in the art (see, e.g., Ausubel et al, eds, 1994, Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York, which is incorporated by reference herein in its entirety). Exemplary immunoassays are described briefly below (but are not intended by way of limitation).

Immunoprecipitation protocols generally comprise lysing a population of cells in a lysis buffer such as RIPA buffer (1% NP-40 or Triton X- 100, 1% sodium deoxycholate, 0.1% SDS, 0.15 M NaCl, 0.01 M sodium phosphate at pH 7.2, 1% Trasylol) supplemented with protein phosphatase and/or protease inhibitors (e.g., EDTA, PMSF, aprotinin, sodium vanadate), adding the antibody of interest to the cell lysate, incubating for a period of time (e.g., 1-4 hours) at 4° C, adding protein A and/or protein G sepharose beads to the cell lysate, incubating for about an hour or more at 4° C, washing the beads in lysis buffer and resuspending the beads in SDS/sample buffer. The ability of the antibody of interest to immunoprecipitate a particular antigen can be assessed by, e.g., western blot analysis. One of skill in the art would be knowledgeable as to the parameters that can be modified to increase the binding of the antibody to an antigen and decrease the background (e.g., pre-clearing the cell lysate with sepharose beads). For further discussion regarding immunoprecipitation protocols see, e.g., Ausubel et al, eds, 1994, Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York at 10.16.1.

Western blot analysis generally comprises preparing protein samples, electrophoresis of the protein samples in a polyacrylamide gel (e.g., 8%- 20% SDS-PAGE depending on the molecular weight of the antigen), transferring the protein sample from the polyacrylamide gel to a membrane such as nitrocellulose, PVDF or nylon, blocking the membrane in blocking solution (e.g., PBS with 3% BSA or non-fat milk), washing the membrane in washing buffer

(e.g., PBS-Tween 20), blocking the membrane with primary antibody (the antibody of interest) diluted in blocking buffer, washing the membrane in washing buffer, blocking the membrane with a secondary antibody (which recognizes the primary antibody, e.g., an anti-human antibody) conjugated to an enzymatic substrate (e.g., horseradish peroxidase or alkaline phosphatase) or radioactive molecule (e.g., ^{32}P or ^{125}I) diluted in blocking buffer, washing the membrane in wash buffer, and detecting the presence of the antigen. One of skill in the art would be knowledgeable as to the parameters that can be modified to increase the signal detected and to reduce the background noise. For further discussion regarding western blot protocols see, e.g., Ausubel et al, eds, 1994, Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York at 10.8.1.

ELISAs comprise preparing antigen, coating the well of a 96 well microtiter plate with the antigen, adding the antibody of interest conjugated to a detectable compound such as an enzymatic substrate (e.g., horseradish peroxidase or alkaline phosphatase) to the well and incubating for a period of time, and detecting the presence of the antigen. In ELISAs the antibody of interest does not have to be conjugated to a detectable compound; instead, a second antibody (which recognizes the antibody of interest) conjugated to a detectable compound may be added to the well. Further, instead of coating the well with the antigen, the antibody may be coated to the well. In this case, a second antibody conjugated to a detectable compound may be added following the addition of the antigen of interest to the coated well. One of skill in the art would be knowledgeable as to the parameters that can be modified to increase the signal detected as well as other variations of ELISAs known in the art. For further discussion regarding ELISAs see, e.g., Ausubel et al, eds, 1994, Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York at 11.2.1.

The binding affinity of an antibody to an antigen and the off-rate of an antibody-antigen interaction can be determined by competitive binding assays. One example of a competitive binding assay is a radioimmunoassay comprising the incubation of labeled antigen (e.g., ^3H or ^{125}I) with the antibody of interest in the presence of increasing amounts of unlabeled antigen, and the detection of the antibody bound to the labeled antigen. The affinity of the antibody of interest for a particular antigen and the binding off-rates can be determined from the data by scatchard plot analysis. Competition with a second antibody can also be determined using radioimmunoassays. In this case, the antigen is incubated with

antibody of interest conjugated to a labeled compound (e.g., ^3H or ^{125}I) in the presence of increasing amounts of an unlabeled second antibody.

Therapeutic Uses

5 The present invention is further directed to antibody-based therapies which involve administering antibodies of the invention to an animal, preferably a mammal, and most preferably a human, patient for treating one or more of the disclosed diseases, disorders, or conditions. Therapeutic compounds of the invention include, but are not limited to, antibodies of the invention (including fragments, analogs and derivatives thereof as described
10 herein) and nucleic acids encoding antibodies of the invention (including fragments, analogs and derivatives thereof and anti-idiotypic antibodies as described herein). The antibodies of the invention can be used to treat, inhibit or prevent diseases, disorders or conditions associated with aberrant expression and/or activity of a polypeptide of the invention, including, but not limited to, any one or more of the diseases, disorders, or conditions
15 described herein. The treatment and/or prevention of diseases, disorders, or conditions associated with aberrant expression and/or activity of a polypeptide of the invention includes, but is not limited to, alleviating symptoms associated with those diseases, disorders or conditions. Antibodies of the invention may be provided in pharmaceutically acceptable compositions as known in the art or as described herein.

20 A summary of the ways in which the antibodies of the present invention may be used therapeutically includes binding polynucleotides or polypeptides of the present invention locally or systemically in the body or by direct cytotoxicity of the antibody, e.g. as mediated by complement (CDC) or by effector cells (ADCC). Some of these approaches are described in more detail below. Armed with the teachings provided herein, one of ordinary skill in the
25 art will know how to use the antibodies of the present invention for diagnostic, monitoring or therapeutic purposes without undue experimentation.

 The antibodies of this invention may be advantageously utilized in combination with other monoclonal or chimeric antibodies, or with lymphokines or hematopoietic growth factors (such as, e.g., IL-2, IL-3 and IL-7), for example, which serve to increase the number
30 or activity of effector cells which interact with the antibodies.

 The antibodies of the invention may be administered alone or in combination with other types of treatments (e.g., radiation therapy, chemotherapy, hormonal therapy,

immunotherapy and anti-tumor agents). Generally, administration of products of a species origin or species reactivity (in the case of antibodies) that is the same species as that of the patient is preferred. Thus, in a preferred embodiment, human antibodies, fragments derivatives, analogs, or nucleic acids, are administered to a human patient for therapy or prophylaxis.

It is preferred to use high affinity and/or potent in vivo inhibiting and/or neutralizing antibodies against polypeptides or polynucleotides of the present invention, fragments or regions thereof, for both immunoassays directed to and therapy of disorders related to polynucleotides or polypeptides, including fragments thereof, of the present invention. Such antibodies, fragments, or regions, will preferably have an affinity for polynucleotides or polypeptides of the invention, including fragments thereof. Preferred binding affinities include those with a dissociation constant or K_d less than 5×10^{-2} M, 10^{-2} M, 5×10^{-3} M, 10^{-3} M, 5×10^{-4} M, 10^{-4} M, 5×10^{-5} M, 10^{-5} M, 5×10^{-6} M, 10^{-6} M, 5×10^{-7} M, 10^{-7} M, 5×10^{-8} M, 10^{-8} M, 5×10^{-9} M, 10^{-9} M, 5×10^{-10} M, 10^{-10} M, 5×10^{-11} M, 10^{-11} M, 5×10^{-12} M, 10^{-12} M, 5×10^{-13} M, 10^{-13} M, 5×10^{-14} M, 10^{-14} M, 5×10^{-15} M, and 10^{-15} M.

Gene Therapy

In a specific embodiment, nucleic acids comprising sequences encoding antibodies or functional derivatives thereof, are administered to treat, inhibit or prevent a disease or disorder associated with aberrant expression and/or activity of a polypeptide of the invention, by way of gene therapy. Gene therapy refers to therapy performed by the administration to a subject of an expressed or expressible nucleic acid. In this embodiment of the invention, the nucleic acids produce their encoded protein that mediates a therapeutic effect.

Any of the methods for gene therapy available in the art can be used according to the present invention. Exemplary methods are described below.

For general reviews of the methods of gene therapy, see Goldspiel et al., *Clinical Pharmacy* 12:488-505 (1993); Wu and Wu, *Biotherapy* 3:87-95 (1991); Tolstoshev, *Ann. Rev. Pharmacol. Toxicol.* 32:573-596 (1993); Mulligan, *Science* 260:926-932 (1993); and Morgan and Anderson, *Ann. Rev. Biochem.* 62:191-217 (1993); May, *TIBTECH* 11(5):155-215 (1993). Methods commonly known in the art of recombinant DNA technology which can be used are described in Ausubel et al. (eds.), *Current Protocols in Molecular Biology*, John

Wiley & Sons, NY (1993); and Kriegler, Gene Transfer and Expression, A Laboratory Manual, Stockton Press, NY (1990).

In a preferred aspect, the compound comprises nucleic acid sequences encoding an antibody, said nucleic acid sequences being part of expression vectors that express the antibody or fragments or chimeric proteins or heavy or light chains thereof in a suitable host. In particular, such nucleic acid sequences have promoters operably linked to the antibody coding region, said promoter being inducible or constitutive, and, optionally, tissue-specific. In another particular embodiment, nucleic acid molecules are used in which the antibody coding sequences and any other desired sequences are flanked by regions that promote homologous recombination at a desired site in the genome, thus providing for intrachromosomal expression of the antibody encoding nucleic acids (Koller and Smithies, Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); Zijlstra et al., Nature 342:435-438 (1989). In specific embodiments, the expressed antibody molecule is a single chain antibody; alternatively, the nucleic acid sequences include sequences encoding both the heavy and light chains, or fragments thereof, of the antibody.

Delivery of the nucleic acids into a patient may be either direct, in which case the patient is directly exposed to the nucleic acid or nucleic acid-carrying vectors, or indirect, in which case, cells are first transformed with the nucleic acids in vitro, then transplanted into the patient. These two approaches are known, respectively, as in vivo or ex vivo gene therapy.

In a specific embodiment, the nucleic acid sequences are directly administered in vivo, where it is expressed to produce the encoded product. This can be accomplished by any of numerous methods known in the art, e.g., by constructing them as part of an appropriate nucleic acid expression vector and administering it so that they become intracellular, e.g., by infection using defective or attenuated retrovirals or other viral vectors (see U.S. Patent No. 4,980,286), or by direct injection of naked DNA, or by use of microparticle bombardment (e.g., a gene gun; Biolistic, Dupont), or coating with lipids or cell-surface receptors or transfecting agents, encapsulation in liposomes, microparticles, or microcapsules, or by administering them in linkage to a peptide which is known to enter the nucleus, by administering it in linkage to a ligand subject to receptor-mediated endocytosis (see, e.g., Wu and Wu, J. Biol. Chem. 262:4429-4432 (1987)) (which can be used to target cell types specifically expressing the receptors), etc. In another embodiment, nucleic acid-

ligand complexes can be formed in which the ligand comprises a fusogenic viral peptide to disrupt endosomes, allowing the nucleic acid to avoid lysosomal degradation. In yet another embodiment, the nucleic acid can be targeted in vivo for cell specific uptake and expression, by targeting a specific receptor (see, e.g., PCT Publications WO 92/06180; WO 92/22635; WO92/20316; WO93/14188, WO 93/20221). Alternatively, the nucleic acid can be introduced intracellularly and incorporated within host cell DNA for expression, by homologous recombination (Koller and Smithies, Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); Zijlstra et al., Nature 342:435-438 (1989)).

In a specific embodiment, viral vectors that contains nucleic acid sequences encoding an antibody of the invention are used. For example, a retroviral vector can be used (see Miller et al., Meth. Enzymol. 217:581-599 (1993)). These retroviral vectors contain the components necessary for the correct packaging of the viral genome and integration into the host cell DNA. The nucleic acid sequences encoding the antibody to be used in gene therapy are cloned into one or more vectors, which facilitates delivery of the gene into a patient. More detail about retroviral vectors can be found in Boesen et al., Biotherapy 6:291-302 (1994), which describes the use of a retroviral vector to deliver the *mdr1* gene to hematopoietic stem cells in order to make the stem cells more resistant to chemotherapy. Other references illustrating the use of retroviral vectors in gene therapy are: Clowes et al., J. Clin. Invest. 93:644-651 (1994); Kiem et al., Blood 83:1467-1473 (1994); Salmons and Gunzberg, Human Gene Therapy 4:129-141 (1993); and Grossman and Wilson, Curr. Opin. in Genetics and Devel. 3:110-114 (1993).

Adenoviruses are other viral vectors that can be used in gene therapy. Adenoviruses are especially attractive vehicles for delivering genes to respiratory epithelia. Adenoviruses naturally infect respiratory epithelia where they cause a mild disease. Other targets for adenovirus-based delivery systems are liver, the central nervous system, endothelial cells, and muscle. Adenoviruses have the advantage of being capable of infecting non-dividing cells. Kozarsky and Wilson, Current Opinion in Genetics and Development 3:499-503 (1993) present a review of adenovirus-based gene therapy. Bout et al., Human Gene Therapy 5:3-10 (1994) demonstrated the use of adenovirus vectors to transfer genes to the respiratory epithelia of rhesus monkeys. Other instances of the use of adenoviruses in gene therapy can be found in Rosenfeld et al., Science 252:431-434 (1991); Rosenfeld et al., Cell 68:143-155 (1992); Mastrangeli et al., J. Clin. Invest. 91:225-234 (1993); PCT Publication

WO94/12649; and Wang, et al., Gene Therapy 2:775-783 (1995). In a preferred embodiment, adenovirus vectors are used.

Adeno-associated virus (AAV) has also been proposed for use in gene therapy (Walsh et al., Proc. Soc. Exp. Biol. Med. 204:289-300 (1993); U.S. Patent No. 5,436,146).

5 Another approach to gene therapy involves transferring a gene to cells in tissue culture by such methods as electroporation, lipofection, calcium phosphate mediated transfection, or viral infection. Usually, the method of transfer includes the transfer of a selectable marker to the cells. The cells are then placed under selection to isolate those cells that have taken up and are expressing the transferred gene. Those cells are then delivered to a
10 patient.

In this embodiment, the nucleic acid is introduced into a cell prior to administration in vivo of the resulting recombinant cell. Such introduction can be carried out by any method known in the art, including but not limited to transfection, electroporation, microinjection, infection with a viral or bacteriophage vector containing the nucleic acid sequences, cell
15 fusion, chromosome-mediated gene transfer, microcell-mediated gene transfer, spheroplast fusion, etc. Numerous techniques are known in the art for the introduction of foreign genes into cells (see, e.g., Loeffler and Behr, Meth. Enzymol. 217:599-618 (1993); Cohen et al., Meth. Enzymol. 217:618-644 (1993); Cline, Pharmac. Ther. 29:69-92m (1985) and may be used in accordance with the present invention, provided that the necessary developmental
20 and physiological functions of the recipient cells are not disrupted. The technique should provide for the stable transfer of the nucleic acid to the cell, so that the nucleic acid is expressible by the cell and preferably heritable and expressible by its cell progeny.

The resulting recombinant cells can be delivered to a patient by various methods known in the art. Recombinant blood cells (e.g., hematopoietic stem or progenitor cells) are
25 preferably administered intravenously. The amount of cells envisioned for use depends on the desired effect, patient state, etc., and can be determined by one skilled in the art.

Cells into which a nucleic acid can be introduced for purposes of gene therapy encompass any desired, available cell type, and include but are not limited to epithelial cells, endothelial cells, keratinocytes, fibroblasts, muscle cells, hepatocytes; blood cells such as
30 Tlymphocytes, Blymphocytes, monocytes, macrophages, neutrophils, eosinophils, megakaryocytes, granulocytes; various stem or progenitor cells, in particular hematopoietic

stem or progenitor cells, e.g., as obtained from bone marrow, umbilical cord blood, peripheral blood, fetal liver, etc.

In a preferred embodiment, the cell used for gene therapy is autologous to the patient.

In an embodiment in which recombinant cells are used in gene therapy, nucleic acid sequences encoding an antibody are introduced into the cells such that they are expressible by the cells or their progeny, and the recombinant cells are then administered in vivo for therapeutic effect. In a specific embodiment, stem or progenitor cells are used. Any stem and/or progenitor cells which can be isolated and maintained in vitro can potentially be used in accordance with this embodiment of the present invention (see e.g. PCT Publication WO 94/08598; Stemple and Anderson, Cell 71:973-985 (1992); Rheinwald, Meth. Cell Bio. 21A:229 (1980); and Pittelkow and Scott, Mayo Clinic Proc. 61:771 (1986)).

In a specific embodiment, the nucleic acid to be introduced for purposes of gene therapy comprises an inducible promoter operably linked to the coding region, such that expression of the nucleic acid is controllable by controlling the presence or absence of the appropriate inducer of transcription. Demonstration of Therapeutic or Prophylactic Activity

The compounds or pharmaceutical compositions of the invention are preferably tested in vitro, and then in vivo for the desired therapeutic or prophylactic activity, prior to use in humans. For example, in vitro assays to demonstrate the therapeutic or prophylactic utility of a compound or pharmaceutical composition include, the effect of a compound on a cell line or a patient tissue sample. The effect of the compound or composition on the cell line and/or tissue sample can be determined utilizing techniques known to those of skill in the art including, but not limited to, rosette formation assays and cell lysis assays. In accordance with the invention, in vitro assays which can be used to determine whether administration of a specific compound is indicated, include in vitro cell culture assays in which a patient tissue sample is grown in culture, and exposed to or otherwise administered a compound, and the effect of such compound upon the tissue sample is observed.

Therapeutic/Prophylactic Administration and Composition

The invention provides methods of treatment, inhibition and prophylaxis by administration to a subject of an effective amount of a compound or pharmaceutical composition of the invention, preferably a polypeptide or antibody of the invention. In a preferred aspect, the compound is substantially purified (e.g., substantially free from

substances that limit its effect or produce undesired side-effects). The subject is preferably an animal, including but not limited to animals such as cows, pigs, horses, chickens, cats, dogs, etc., and is preferably a mammal, and most preferably human.

Formulations and methods of administration that can be employed when the compound comprises a nucleic acid or an immunoglobulin are described above; additional appropriate formulations and routes of administration can be selected from among those described herein below.

Various delivery systems are known and can be used to administer a compound of the invention, e.g., encapsulation in liposomes, microparticles, microcapsules, recombinant cells capable of expressing the compound, receptor-mediated endocytosis (see, e.g., Wu and Wu, J. Biol. Chem. 262:4429-4432 (1987)), construction of a nucleic acid as part of a retroviral or other vector, etc. Methods of introduction include but are not limited to intradermal, intramuscular, intraperitoneal, intravenous, subcutaneous, intranasal, epidural, and oral routes. The compounds or compositions may be administered by any convenient route, for example by infusion or bolus injection, by absorption through epithelial or mucocutaneous linings (e.g., oral mucosa, rectal and intestinal mucosa, etc.) and may be administered together with other biologically active agents. Administration can be systemic or local. In addition, it may be desirable to introduce the pharmaceutical compounds or compositions of the invention into the central nervous system by any suitable route, including intraventricular and intrathecal injection; intraventricular injection may be facilitated by an intraventricular catheter, for example, attached to a reservoir, such as an Ommaya reservoir. Pulmonary administration can also be employed, e.g., by use of an inhaler or nebulizer, and formulation with an aerosolizing agent.

In a specific embodiment, it may be desirable to administer the pharmaceutical compounds or compositions of the invention locally to the area in need of treatment; this may be achieved by, for example, and not by way of limitation, local infusion during surgery, topical application, e.g., in conjunction with a wound dressing after surgery, by injection, by means of a catheter, by means of a suppository, or by means of an implant, said implant being of a porous, non-porous, or gelatinous material, including membranes, such as sialastic membranes, or fibers. Preferably, when administering a protein, including an antibody, of the invention, care must be taken to use materials to which the protein does not absorb.

In another embodiment, the compound or composition can be delivered in a vesicle, in particular a liposome (see Langer, *Science* 249:1527-1533 (1990); Treat et al., in *Liposomes in the Therapy of Infectious Disease and Cancer*, Lopez-Berestein and Fidler (eds.), Liss, New York, pp. 353- 365 (1989); Lopez-Berestein, *ibid.*, pp. 317-327; see generally *ibid.*)

In yet another embodiment, the compound or composition can be delivered in a controlled release system. In one embodiment, a pump may be used (see Langer, *supra*; Sefton, *CRC Crit. Ref. Biomed. Eng.* 14:201 (1987); Buchwald et al., *Surgery* 88:507 (1980); Saudek et al., *N. Engl. J. Med.* 321:574 (1989)). In another embodiment, polymeric materials can be used (see *Medical Applications of Controlled Release*, Langer and Wise (eds.), CRC Pres., Boca Raton, Florida (1974); *Controlled Drug Bioavailability, Drug Product Design and Performance*, Smolen and Ball (eds.), Wiley, New York (1984); Ranger and Peppas, J., *Macromol. Sci. Rev. Macromol. Chem.* 23:61 (1983); see also Levy et al., *Science* 228:190 (1985); During et al., *Ann. Neurol.* 25:351 (1989); Howard et al., *J. Neurosurg.* 71:105 (1989)). In yet another embodiment, a controlled release system can be placed in proximity of the therapeutic target, i.e., the brain, thus requiring only a fraction of the systemic dose (see, e.g., Goodson, in *Medical Applications of Controlled Release*, *supra*, vol. 2, pp. 115-138 (1984)).

Other controlled release systems are discussed in the review by Langer (*Science* 249:1527-1533 (1990)).

In a specific embodiment where the compound of the invention is a nucleic acid encoding a protein, the nucleic acid can be administered *in vivo* to promote expression of its encoded protein, by constructing it as part of an appropriate nucleic acid expression vector and administering it so that it becomes intracellular, e.g., by use of a retroviral vector (see U.S. Patent No. 4,980,286), or by direct injection, or by use of microparticle bombardment (e.g., a gene gun; Biolistic, Dupont), or coating with lipids or cell-surface receptors or transfecting agents, or by administering it in linkage to a homeobox- like peptide which is known to enter the nucleus (see e.g., Joliot et al., *Proc. Natl. Acad. Sci. USA* 88:1864-1868 (1991)), etc. Alternatively, a nucleic acid can be introduced intracellularly and incorporated within host cell DNA for expression, by homologous recombination.

The present invention also provides pharmaceutical compositions. Such compositions comprise a therapeutically effective amount of a compound, and a pharmaceutically

acceptable carrier. In a specific embodiment, the term "pharmaceutically acceptable" means approved by a regulatory agency of the Federal or a state government or listed in the U.S. Pharmacopeia or other generally recognized pharmacopeia for use in animals, and more particularly in humans. The term "carrier" refers to a diluent, adjuvant, excipient, or vehicle with which the therapeutic is administered. Such pharmaceutical carriers can be sterile liquids, such as water and oils, including those of petroleum, animal, vegetable or synthetic origin, such as peanut oil, soybean oil, mineral oil, sesame oil and the like. Water is a preferred carrier when the pharmaceutical composition is administered intravenously. Saline solutions and aqueous dextrose and glycerol solutions can also be employed as liquid carriers, particularly for injectable solutions. Suitable pharmaceutical excipients include starch, glucose, lactose, sucrose, gelatin, malt, rice, flour, chalk, silica gel, sodium stearate, glycerol monostearate, talc, sodium chloride, dried skim milk, glycerol, propylene, glycol, water, ethanol and the like. The composition, if desired, can also contain minor amounts of wetting or emulsifying agents, or pH buffering agents. These compositions can take the form of solutions, suspensions, emulsion, tablets, pills, capsules, powders, sustained-release formulations and the like. The composition can be formulated as a suppository, with traditional binders and carriers such as triglycerides. Oral formulation can include standard carriers such as pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharine, cellulose, magnesium carbonate, etc. Examples of suitable pharmaceutical carriers are described in "Remington's Pharmaceutical Sciences" by E.W. Martin. Such compositions will contain a therapeutically effective amount of the compound, preferably in purified form, together with a suitable amount of carrier so as to provide the form for proper administration to the patient. The formulation should suit the mode of administration.

In a preferred embodiment, the composition is formulated in accordance with routine procedures as a pharmaceutical composition adapted for intravenous administration to human beings. Typically, compositions for intravenous administration are solutions in sterile isotonic aqueous buffer. Where necessary, the composition may also include a solubilizing agent and a local anesthetic such as lignocaine to ease pain at the site of the injection. Generally, the ingredients are supplied either separately or mixed together in unit dosage form, for example, as a dry lyophilized powder or water free concentrate in a hermetically sealed container such as an ampoule or sachette indicating the quantity of active agent.

Where the composition is to be administered by infusion, it can be dispensed with an infusion bottle containing sterile pharmaceutical grade water or saline. Where the composition is administered by injection, an ampoule of sterile water for injection or saline can be provided so that the ingredients may be mixed prior to administration.

5 The compounds of the invention can be formulated as neutral or salt forms. Pharmaceutically acceptable salts include those formed with anions such as those derived from hydrochloric, phosphoric, acetic, oxalic, tartaric acids, etc., and those formed with cations such as those derived from sodium, potassium, ammonium, calcium, ferric hydroxides, isopropylamine, triethylamine, 2-ethylamino ethanol, histidine, procaine, etc.

10 The amount of the compound of the invention which will be effective in the treatment, inhibition and prevention of a disease or disorder associated with aberrant expression and/or activity of a polypeptide of the invention can be determined by standard clinical techniques. In addition, in vitro assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also depend
15 on the route of administration, and the seriousness of the disease or disorder, and should be decided according to the judgment of the practitioner and each patient's circumstances. Effective doses may be extrapolated from dose-response curves derived from in vitro or animal model test systems.

For antibodies, the dosage administered to a patient is typically 0.1 mg/kg to 100
20 mg/kg of the patient's body weight. Preferably, the dosage administered to a patient is between 0.1 mg/kg and 20 mg/kg of the patient's body weight, more preferably 1 mg/kg to 10 mg/kg of the patient's body weight. Generally, human antibodies have a longer half-life within the human body than antibodies from other species due to the immune response to the foreign polypeptides. Thus, lower dosages of human antibodies and less frequent
25 administration is often possible. Further, the dosage and frequency of administration of antibodies of the invention may be reduced by enhancing uptake and tissue penetration (e.g., into the brain) of the antibodies by modifications such as, for example, lipidation.

The invention also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the pharmaceutical compositions of
30 the invention. Optionally associated with such container(s) can be a notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of

pharmaceuticals or biological products, which notice reflects approval by the agency of manufacture, use or sale for human administration.

Diagnosis and Imaging

5 Labeled antibodies, and derivatives and analogs thereof, which specifically bind to a polypeptide of interest can be used for diagnostic purposes to detect, diagnose, or monitor diseases, disorders, and/or conditions associated with the aberrant expression and/or activity of a polypeptide of the invention. The invention provides for the detection of aberrant expression of a polypeptide of interest, comprising (a) assaying the expression of the
10 polypeptide of interest in cells or body fluid of an individual using one or more antibodies specific to the polypeptide interest and (b) comparing the level of gene expression with a standard gene expression level, whereby an increase or decrease in the assayed polypeptide gene expression level compared to the standard expression level is indicative of aberrant expression.

15 The invention provides a diagnostic assay for diagnosing a disorder, comprising (a) assaying the expression of the polypeptide of interest in cells or body fluid of an individual using one or more antibodies specific to the polypeptide interest and (b) comparing the level of gene expression with a standard gene expression level, whereby an increase or decrease in the assayed polypeptide gene expression level compared to the standard expression level is
20 indicative of a particular disorder. With respect to cancer, the presence of a relatively high amount of transcript in biopsied tissue from an individual may indicate a predisposition for the development of the disease, or may provide a means for detecting the disease prior to the appearance of actual clinical symptoms. A more definitive diagnosis of this type may allow health professionals to employ preventative measures or aggressive treatment earlier thereby
25 preventing the development or further progression of the cancer.

Antibodies of the invention can be used to assay protein levels in a biological sample using classical immunohistological methods known to those of skill in the art (e.g., see Jalkanen, et al., J. Cell. Biol. 101:976-985 (1985); Jalkanen, et al., J. Cell . Biol. 105:3087-3096 (1987)). Other antibody-based methods useful for detecting protein gene expression
30 include immunoassays, such as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). Suitable antibody assay labels are known in the art and include enzyme labels, such as, glucose oxidase; radioisotopes, such as iodine (125I, 121I), carbon

(14C), sulfur (35S), tritium (3H), indium (112In), and technetium (99Tc); luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

One aspect of the invention is the detection and diagnosis of a disease or disorder associated with aberrant expression of a polypeptide of interest in an animal, preferably a mammal and most preferably a human. In one embodiment, diagnosis comprises: a) administering (for example, parenterally, subcutaneously, or intraperitoneally) to a subject an effective amount of a labeled molecule which specifically binds to the polypeptide of interest; b) waiting for a time interval following the administering for permitting the labeled molecule to preferentially concentrate at sites in the subject where the polypeptide is expressed (and for unbound labeled molecule to be cleared to background level); c) determining background level; and d) detecting the labeled molecule in the subject, such that detection of labeled molecule above the background level indicates that the subject has a particular disease or disorder associated with aberrant expression of the polypeptide of interest. Background level can be determined by various methods including, comparing the amount of labeled molecule detected to a standard value previously determined for a particular system.

It will be understood in the art that the size of the subject and the imaging system used will determine the quantity of imaging moiety needed to produce diagnostic images. In the case of a radioisotope moiety, for a human subject, the quantity of radioactivity injected will normally range from about 5 to 20 millicuries of 99mTc. The labeled antibody or antibody fragment will then preferentially accumulate at the location of cells which contain the specific protein. In vivo tumor imaging is described in S.W. Burchiel et al., "Immunopharmacokinetics of Radiolabeled Antibodies and Their Fragments." (Chapter 13 in Tumor Imaging: The Radiochemical Detection of Cancer, S.W. Burchiel and B. A. Rhodes, eds., Masson Publishing Inc. (1982).

Depending on several variables, including the type of label used and the mode of administration, the time interval following the administration for permitting the labeled molecule to preferentially concentrate at sites in the subject and for unbound labeled molecule to be cleared to background level is 6 to 48 hours or 6 to 24 hours or 6 to 12 hours. In another embodiment the time interval following administration is 5 to 20 days or 5 to 10 days.

In an embodiment, monitoring of the disease or disorder is carried out by repeating the method for diagnosing the disease or disease, for example, one month after initial diagnosis, six months after initial diagnosis, one year after initial diagnosis, etc.

Presence of the labeled molecule can be detected in the patient using methods known in the art for in vivo scanning. These methods depend upon the type of label used. Skilled artisans will be able to determine the appropriate method for detecting a particular label. Methods and devices that may be used in the diagnostic methods of the invention include, but are not limited to, computed tomography (CT), whole body scan such as position emission tomography (PET), magnetic resonance imaging (MRI), and sonography.

In a specific embodiment, the molecule is labeled with a radioisotope and is detected in the patient using a radiation responsive surgical instrument (Thurston et al., U.S. Patent No. 5,441,050). In another embodiment, the molecule is labeled with a fluorescent compound and is detected in the patient using a fluorescence responsive scanning instrument. In another embodiment, the molecule is labeled with a positron emitting metal and is detected in the patient using positron emission-tomography. In yet another embodiment, the molecule is labeled with a paramagnetic label and is detected in a patient using magnetic resonance imaging (MRI).

Kits

The present invention provides kits that can be used in the above methods. In one embodiment, a kit comprises an antibody of the invention, preferably a purified antibody, in one or more containers. In a specific embodiment, the kits of the present invention contain a substantially isolated polypeptide comprising an epitope which is specifically immunoreactive with an antibody included in the kit. Preferably, the kits of the present invention further comprise a control antibody which does not react with the polypeptide of interest. In another specific embodiment, the kits of the present invention contain a means for detecting the binding of an antibody to a polypeptide of interest (e.g., the antibody may be conjugated to a detectable substrate such as a fluorescent compound, an enzymatic substrate, a radioactive compound or a luminescent compound, or a second antibody which recognizes the first antibody may be conjugated to a detectable substrate).

In another specific embodiment of the present invention, the kit is a diagnostic kit for use in screening serum containing antibodies specific against proliferative and/or cancerous

polynucleotides and polypeptides. Such a kit may include a control antibody that does not react with the polypeptide of interest. Such a kit may include a substantially isolated polypeptide antigen comprising an epitope which is specifically immunoreactive with at least one anti-polypeptide antigen antibody. Further, such a kit includes means for detecting the binding of said antibody to the antigen (e.g., the antibody may be conjugated to a fluorescent compound such as fluorescein or rhodamine which can be detected by flow cytometry). In specific embodiments, the kit may include a recombinantly produced or chemically synthesized polypeptide antigen. The polypeptide antigen of the kit may also be attached to a solid support.

In a more specific embodiment the detecting means of the above-described kit includes a solid support to which said polypeptide antigen is attached. Such a kit may also include a non-attached reporter-labeled anti-human antibody. In this embodiment, binding of the antibody to the polypeptide antigen can be detected by binding of the said reporter-labeled antibody.

In an additional embodiment, the invention includes a diagnostic kit for use in screening serum containing antigens of the polypeptide of the invention. The diagnostic kit includes a substantially isolated antibody specifically immunoreactive with polypeptide or polynucleotide antigens, and means for detecting the binding of the polynucleotide or polypeptide antigen to the antibody. In one embodiment, the antibody is attached to a solid support. In a specific embodiment, the antibody may be a monoclonal antibody. The detecting means of the kit may include a second, labeled monoclonal antibody. Alternatively, or in addition, the detecting means may include a labeled, competing antigen.

In one diagnostic configuration, test serum is reacted with a solid phase reagent having a surface-bound antigen obtained by the methods of the present invention. After binding with specific antigen antibody to the reagent and removing unbound serum components by washing, the reagent is reacted with reporter-labeled anti-human antibody to bind reporter to the reagent in proportion to the amount of bound anti-antigen antibody on the solid support. The reagent is again washed to remove unbound labeled antibody, and the amount of reporter associated with the reagent is determined. Typically, the reporter is an enzyme which is detected by incubating the solid phase in the presence of a suitable fluorometric, luminescent or colorimetric substrate (Sigma, St. Louis, MO).

The solid surface reagent in the above assay is prepared by known techniques for attaching protein material to solid support material, such as polymeric beads, dip sticks, 96-well plate or filter material. These attachment methods generally include non-specific adsorption of the protein to the support or covalent attachment of the protein, typically through a free amine group, to a chemically reactive group on the solid support, such as an activated carboxyl, hydroxyl, or aldehyde group. Alternatively, streptavidin coated plates can be used in conjunction with biotinylated antigen(s).

Thus, the invention provides an assay system or kit for carrying out this diagnostic method. The kit generally includes a support with surface-bound recombinant antigens, and a reporter-labeled anti-human antibody for detecting surface-bound anti-antigen antibody.

Uses of the Polynucleotides

Each of the polynucleotides identified herein can be used in numerous ways as reagents. The following description should be considered exemplary and utilizes known techniques.

The polynucleotides of the present invention are useful for chromosome identification. There exists an ongoing need to identify new chromosome markers, since few chromosome marking reagents, based on actual sequence data (repeat polymorphisms), are presently available. Each sequence is specifically targeted to and can hybridize with a particular location on an individual human chromosome, thus each polynucleotide of the present invention can routinely be used as a chromosome marker using techniques known in the art.

Briefly, sequences can be mapped to chromosomes by preparing PCR primers (preferably at least 15 bp (e.g., 15-25 bp) from the sequences shown in SEQ ID NO:X. Primers can optionally be selected using computer analysis so that primers do not span more than one predicted exon in the genomic DNA. These primers are then used for PCR screening of somatic cell hybrids containing individual human chromosomes. Only those hybrids containing the human gene corresponding to SEQ ID NO:X will yield an amplified fragment.

Similarly, somatic hybrids provide a rapid method of PCR mapping the polynucleotides to particular chromosomes. Three or more clones can be assigned per day using a single thermal cycler. Moreover, sublocalization of the polynucleotides can be

achieved with panels of specific chromosome fragments. Other gene mapping strategies that can be used include in situ hybridization, prescreening with labeled flow-sorted chromosomes, preselection by hybridization to construct chromosome specific-cDNA libraries, and computer mapping techniques (See, e.g., Shuler, Trends Biotechnol 16:456-459 (1998) which is hereby incorporated by reference in its entirety).

Precise chromosomal location of the polynucleotides can also be achieved using fluorescence in situ hybridization (FISH) of a metaphase chromosomal spread. This technique uses polynucleotides as short as 500 or 600 bases; however, polynucleotides 2,000-4,000 bp are preferred. For a review of this technique, see Verma et al., "Human Chromosomes: a Manual of Basic Techniques," Pergamon Press, New York (1988).

For chromosome mapping, the polynucleotides can be used individually (to mark a single chromosome or a single site on that chromosome) or in panels (for marking multiple sites and/or multiple chromosomes).

Thus, the present invention also provides a method for chromosomal localization which involves (a) preparing PCR primers from the polynucleotide sequences in Table 1 and SEQ ID NO:X and (b) screening somatic cell hybrids containing individual chromosomes.

The polynucleotides of the present invention would likewise be useful for radiation hybrid mapping, HAPPY mapping, and long range restriction mapping. For a review of these techniques and others known in the art, see, e.g., Dear, "Genome Mapping: A Practical Approach," IRL Press at Oxford University Press, London (1997); Aydin, J. Mol. Med. 77:691-694 (1999); Hacia et al., Mol. Psychiatry 3:483-492 (1998); Herrick et al., Chromosome Res. 7:409-423 (1999); Hamilton et al., Methods Cell Biol. 62:265-280 (2000); and/or Ott, J. Hered. 90:68-70 (1999) each of which is hereby incorporated by reference in its entirety.

Once a polynucleotide has been mapped to a precise chromosomal location, the physical position of the polynucleotide can be used in linkage analysis. Linkage analysis establishes coinheritance between a chromosomal location and presentation of a particular disease. (Disease mapping data are found, for example, in V. McKusick, Mendelian Inheritance in Man (available on line through Johns Hopkins University Welch Medical Library).) Assuming 1 megabase mapping resolution and one gene per 20 kb, a cDNA precisely localized to a chromosomal region associated with the disease could be one of 50-500 potential causative genes.

Thus, once coinheritance is established, differences in a polynucleotide of the invention and the corresponding gene between affected and unaffected individuals can be examined. First, visible structural alterations in the chromosomes, such as deletions or translocations, are examined in chromosome spreads or by PCR. If no structural alterations exist, the presence of point mutations are ascertained. Mutations observed in some or all affected individuals, but not in normal individuals, indicates that the mutation may cause the disease. However, complete sequencing of the polypeptide and the corresponding gene from several normal individuals is required to distinguish the mutation from a polymorphism. If a new polymorphism is identified, this polymorphic polypeptide can be used for further linkage analysis.

Furthermore, increased or decreased expression of the gene in affected individuals as compared to unaffected individuals can be assessed using the polynucleotides of the invention. Any of these alterations (altered expression, chromosomal rearrangement, or mutation) can be used as a diagnostic or prognostic marker.

Thus, the invention also provides a diagnostic method useful during diagnosis of a disorder, involving measuring the expression level of polynucleotides of the present invention in cells or body fluid from an individual and comparing the measured gene expression level with a standard level of polynucleotide expression level, whereby an increase or decrease in the gene expression level compared to the standard is indicative of a disorder.

In still another embodiment, the invention includes a kit for analyzing samples for the presence of proliferative and/or cancerous polynucleotides derived from a test subject. In a general embodiment, the kit includes at least one polynucleotide probe containing a nucleotide sequence that will specifically hybridize with a polynucleotide of the invention and a suitable container. In a specific embodiment, the kit includes two polynucleotide probes defining an internal region of the polynucleotide of the invention, where each probe has one strand containing a 31'mer-end internal to the region. In a further embodiment, the probes may be useful as primers for polymerase chain reaction amplification.

Where a diagnosis of a related disorder, including, for example, diagnosis of a tumor, has already been made according to conventional methods, the present invention is useful as a prognostic indicator, whereby patients exhibiting enhanced or depressed polynucleotide of the invention expression will experience a worse clinical outcome relative to patients expressing the gene at a level nearer the standard level.

By "measuring the expression level of polynucleotides of the invention" is intended qualitatively or quantitatively measuring or estimating the level of the polypeptide of the invention or the level of the mRNA encoding the polypeptide of the invention in a first biological sample either directly (e.g., by determining or estimating absolute protein level or mRNA level) or relatively (e.g., by comparing to the polypeptide level or mRNA level in a second biological sample). Preferably, the polypeptide level or mRNA level in the first biological sample is measured or estimated and compared to a standard polypeptide level or mRNA level, the standard being taken from a second biological sample obtained from an individual not having the related disorder or being determined by averaging levels from a population of individuals not having a related disorder. As will be appreciated in the art, once a standard polypeptide level or mRNA level is known, it can be used repeatedly as a standard for comparison.

By "biological sample" is intended any biological sample obtained from an individual, body fluid, cell line, tissue culture, or other source which contains polypeptide of the present invention or the corresponding mRNA. As indicated, biological samples include body fluids (such as semen, lymph, sera, plasma, urine, synovial fluid and spinal fluid) which contain the polypeptide of the present invention, and tissue sources found to express the polypeptide of the present invention. Methods for obtaining tissue biopsies and body fluids from mammals are well known in the art. Where the biological sample is to include mRNA, a tissue biopsy is the preferred source.

The method(s) provided above may preferably be applied in a diagnostic method and/or kits in which polynucleotides and/or polypeptides of the invention are attached to a solid support. In one exemplary method, the support may be a "gene chip" or a "biological chip" as described in US Patents 5,837,832, 5,874,219, and 5,856,174. Further, such a gene chip with polynucleotides of the invention attached may be used to identify polymorphisms between the isolated polynucleotide sequences of the invention, with polynucleotides isolated from a test subject. The knowledge of such polymorphisms (i.e. their location, as well as, their existence) would be beneficial in identifying disease loci for many disorders, such as for example, in neural disorders, immune system disorders, muscular disorders, reproductive disorders, gastrointestinal disorders, pulmonary disorders, cardiovascular disorders, renal disorders, proliferative disorders, and/or cancerous diseases and conditions. Such a method is

described in US Patents 5,858,659 and 5,856,104. The US Patents referenced supra are hereby incorporated by reference in their entirety herein.

The present invention encompasses polynucleotides of the present invention that are chemically synthesized, or reproduced as peptide nucleic acids (PNA), or according to other methods known in the art. The use of PNAs would serve as the preferred form if the polynucleotides of the invention are incorporated onto a solid support, or gene chip. For the purposes of the present invention, a peptide nucleic acid (PNA) is a polyamide type of DNA analog and the monomeric units for adenine, guanine, thymine and cytosine are available commercially (Perceptive Biosystems). Certain components of DNA, such as phosphorus, phosphorus oxides, or deoxyribose derivatives, are not present in PNAs. As disclosed by P. E. Nielsen, M. Egholm, R. H. Berg and O. Buchardt, *Science* 254, 1497 (1991); and M. Egholm, O. Buchardt, L. Christensen, C. Behrens, S. M. Freier, D. A. Driver, R. H. Berg, S. K. Kim, B. Norden, and P. E. Nielsen, *Nature* 365, 666 (1993), PNAs bind specifically and tightly to complementary DNA strands and are not degraded by nucleases. In fact, PNA binds more strongly to DNA than DNA itself does. This is probably because there is no electrostatic repulsion between the two strands, and also the polyamide backbone is more flexible. Because of this, PNA/DNA duplexes bind under a wider range of stringency conditions than DNA/DNA duplexes, making it easier to perform multiplex hybridization. Smaller probes can be used than with DNA due to the strong binding. In addition, it is more likely that single base mismatches can be determined with PNA/DNA hybridization because a single mismatch in a PNA/DNA 15-mer lowers the melting point ($T_{sub.m}$) by 8°-20° C, vs. 4°-16° C for the DNA/DNA 15-mer duplex. Also, the absence of charge groups in PNA means that hybridization can be done at low ionic strengths and reduce possible interference by salt during the analysis.

The present invention have uses which include, but are not limited to, detecting cancer in mammals. In particular the invention is useful during diagnosis of pathological cell proliferative neoplasias which include, but are not limited to: acute myelogenous leukemias including acute monocytic leukemia, acute myeloblastic leukemia, acute promyelocytic leukemia, acute myelomonocytic leukemia, acute erythroleukemia, acute megakaryocytic leukemia, and acute undifferentiated leukemia, etc.; and chronic myelogenous leukemias including chronic myelomonocytic leukemia, chronic granulocytic leukemia, etc. Preferred

mammals include monkeys, apes, cats, dogs, cows, pigs, horses, rabbits and humans. Particularly preferred are humans.

Pathological cell proliferative disorders are often associated with inappropriate activation of proto-oncogenes. (Germann, E. P. et al., "The Etiology of Acute Leukemia: Molecular Genetics and Viral Oncology," in Neoplastic Diseases of the Blood, Vol 1., Wiernik, P. H. et al. eds., 161-182 (1985)). Neoplasias are now believed to result from the qualitative alteration of a normal cellular gene product, or from the quantitative modification of gene expression by insertion into the chromosome of a viral sequence, by chromosomal translocation of a gene to a more actively transcribed region, or by some other mechanism.

(Germann et al., supra) It is likely that mutated or altered expression of specific genes is involved in the pathogenesis of some leukemias, among other tissues and cell types. (Germann et al., supra) Indeed, the human counterparts of the oncogenes involved in some animal neoplasias have been amplified or translocated in some cases of human leukemia and carcinoma. (Germann et al., supra)

For example, c-myc expression is highly amplified in the non-lymphocytic leukemia cell line HL-60. When HL-60 cells are chemically induced to stop proliferation, the level of c-myc is found to be downregulated. (International Publication Number WO 91/15580). However, it has been shown that exposure of HL-60 cells to a DNA construct that is complementary to the 5' end of c-myc or c-myb blocks translation of the corresponding mRNAs which downregulates expression of the c-myc or c-myb proteins and causes arrest of cell proliferation and differentiation of the treated cells. (International Publication Number WO 91/15580; Wickstrom et al., Proc. Natl. Acad. Sci. 85:1028 (1988); Anfossi et al., Proc. Natl. Acad. Sci. 86:3379 (1989)). However, the skilled artisan would appreciate the present invention's usefulness is not be limited to treatment of proliferative disorders of hematopoietic cells and tissues, in light of the numerous cells and cell types of varying origins which are known to exhibit proliferative phenotypes.

In addition to the foregoing, a polynucleotide of the present invention can be used to control gene expression through triple helix formation or through antisense DNA or RNA. Antisense techniques are discussed, for example, in Okano, J. Neurochem. 56: 560 (1991); "Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988). Triple helix formation is discussed in, for instance Lee et al., Nucleic Acids Research 6: 3073 (1979); Cooney et al., Science 241: 456 (1988); and Dervan et al., Science

251: 1360 (1991). Both methods rely on binding of the polynucleotide to a complementary DNA or RNA. For these techniques, preferred polynucleotides are usually oligonucleotides 20 to 40 bases in length and complementary to either the region of the gene involved in transcription (triple helix - see Lee et al., Nucl. Acids Res. 6:3073 (1979); Cooney et al.,
5 Science 241:456 (1988); and Dervan et al., Science 251:1360 (1991)) or to the mRNA itself (antisense - Okano, J. Neurochem. 56:560 (1991); Oligodeoxy-nucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988).) Triple helix formation optimally results in a shut-off of RNA transcription from DNA, while antisense RNA hybridization blocks translation of an mRNA molecule into polypeptide. The oligonucleotide
10 described above can also be delivered to cells such that the antisense RNA or DNA may be expressed in vivo to inhibit production of polypeptide of the present invention antigens. Both techniques are effective in model systems, and the information disclosed herein can be used to design antisense or triple helix polynucleotides in an effort to treat disease, and in particular, for the treatment of proliferative diseases and/or conditions.

15 Polynucleotides of the present invention are also useful in gene therapy. One goal of gene therapy is to insert a normal gene into an organism having a defective gene, in an effort to correct the genetic defect. The polynucleotides disclosed in the present invention offer a means of targeting such genetic defects in a highly accurate manner. Another goal is to insert a new gene that was not present in the host genome, thereby producing a new trait in the host
20 cell.

The polynucleotides are also useful for identifying individuals from minute biological samples. The United States military, for example, is considering the use of restriction fragment length polymorphism (RFLP) for identification of its personnel. In this technique, an individual's genomic DNA is digested with one or more restriction enzymes, and probed
25 on a Southern blot to yield unique bands for identifying personnel. This method does not suffer from the current limitations of "Dog Tags" which can be lost, switched, or stolen, making positive identification difficult. The polynucleotides of the present invention can be used as additional DNA markers for RFLP.

The polynucleotides of the present invention can also be used as an alternative to
30 RFLP, by determining the actual base-by-base DNA sequence of selected portions of an individual's genome. These sequences can be used to prepare PCR primers for amplifying and isolating such selected DNA, which can then be sequenced. Using this technique,

individuals can be identified because each individual will have a unique set of DNA sequences. Once an unique ID database is established for an individual, positive identification of that individual, living or dead, can be made from extremely small tissue samples.

5 Forensic biology also benefits from using DNA-based identification techniques as disclosed herein. DNA sequences taken from very small biological samples such as tissues, e.g., hair or skin, or body fluids, e.g., blood, saliva, semen, synovial fluid, amniotic fluid, breast milk, lymph, pulmonary sputum or surfactant, urine, fecal matter, etc., can be amplified using PCR. In one prior art technique, gene sequences amplified from
10 polymorphic loci, such as DQa class II HLA gene, are used in forensic biology to identify individuals. (Erlich, H., PCR Technology, Freeman and Co. (1992).) Once these specific polymorphic loci are amplified, they are digested with one or more restriction enzymes, yielding an identifying set of bands on a Southern blot probed with DNA corresponding to the DQa class II HLA gene. Similarly, polynucleotides of the present invention can be used
15 as polymorphic markers for forensic purposes.

There is also a need for reagents capable of identifying the source of a particular tissue. Such need arises, for example, in forensics when presented with tissue of unknown origin. Appropriate reagents can comprise, for example, DNA probes or primers prepared from the sequences of the present invention, specific to tissues, including but not limited to,
20 prostate, or those listed above for each SEQ ID NO:X. Panels of such reagents can identify tissue by species and/or by organ type. In a similar fashion, these reagents can be used to screen tissue cultures for contamination.

The polynucleotides of the present invention are also useful as hybridization probes for differential identification of the tissue(s) or cell type(s) present in a biological sample.
25 Similarly, polypeptides and antibodies directed to polypeptides of the present invention are useful to provide immunological probes for differential identification of the tissue(s) (e.g., immunohistochemistry assays) or cell type(s) (e.g., immunocytochemistry assays). In addition, for a number of disorders of the above tissues or cells, significantly higher or lower levels of gene expression of the polynucleotides/polypeptides of the present invention may be
30 detected in certain tissues (e.g., tissues expressing polypeptides and/or polynucleotides of the present invention, for example, prostate tissue, and/or cancerous and/or wounded tissues) or bodily fluids (e.g., semen, lymph, serum, plasma, urine, vaginal pool, synovial fluid or spinal

fluid) taken from an individual having such a disorder, relative to a "standard" gene expression level, i.e., the expression level in healthy tissue from an individual not having the disorder.

Thus, the invention provides a diagnostic method of a disorder, which involves: (a) assaying gene expression level in cells or body fluid of an individual; (b) comparing the gene expression level with a standard gene expression level, whereby an increase or decrease in the assayed gene expression level compared to the standard expression level is indicative of a disorder.

In the very least, the polynucleotides of the present invention can be used as molecular weight markers on Southern gels, as diagnostic probes for the presence of a specific mRNA in a particular cell type, as a probe to "subtract-out" known sequences in the process of discovering novel polynucleotides, for selecting and making oligomers for attachment to a "gene chip" or other support, to raise anti-DNA antibodies using DNA immunization techniques, and as an antigen to elicit an immune response.

Uses of the Polypeptides

Each of the polypeptides identified herein can be used in numerous ways. The following description should be considered exemplary and utilizes known techniques.

Polypeptides and antibodies directed to polypeptides of the present invention are useful to provide immunological probes for differential identification of the tissue(s) (e.g., immunohistochemistry assays such as, for example, ABC immunoperoxidase (Hsu et al., J. Histochem. Cytochem. 29:577-580 (1981)) or cell type(s) (e.g., immunocytochemistry assays).

Antibodies can be used to assay levels of polypeptides encoded by polynucleotides of the invention in a biological sample using classical immunohistological methods known to those of skill in the art (e.g., see Jalkanen, et al., J. Cell. Biol. 101:976-985 (1985); Jalkanen, et al., J. Cell. Biol. 105:3087-3096 (1987)). Other antibody-based methods useful for detecting protein gene expression include immunoassays, such as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). Suitable antibody assay labels are known in the art and include enzyme labels, such as, glucose oxidase; radioisotopes, such as iodine (^{131}I , ^{125}I , ^{123}I , ^{121}I), carbon (^{14}C), sulfur (^{35}S), tritium (^3H), indium ($^{115\text{m}}\text{In}$, $^{113\text{m}}\text{In}$, ^{112}In , ^{111}In), and technetium (^{99}Tc , $^{99\text{m}}\text{Tc}$), thallium (^{201}Tl), gallium

(⁶⁸Ga, ⁶⁷Ga), palladium (¹⁰³Pd), molybdenum (⁹⁹Mo), xenon (¹³³Xe), fluorine (¹⁸F), ¹⁵³Sm, ¹⁷⁷Lu, ¹⁵⁹Gd, ¹⁴⁹Pm, ¹⁴⁰La, ¹⁷⁵Yb, ¹⁶⁶Ho, ⁹⁰Y, ⁴⁷Sc, ¹⁸⁶Re, ¹⁸⁸Re, ¹⁴²Pr, ¹⁰⁵Rh, ⁹⁷Ru; luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

5 In addition to assaying levels of polypeptide of the present invention in a biological sample, proteins can also be detected in vivo by imaging. Antibody labels or markers for in vivo imaging of protein include those detectable by X-radiography, NMR or ESR. For X-radiography, suitable labels include radioisotopes such as barium or cesium, which emit detectable radiation but are not overtly harmful to the subject. Suitable markers for NMR and
10 ESR include those with a detectable characteristic spin, such as deuterium, which may be incorporated into the antibody by labeling of nutrients for the relevant hybridoma.

A protein-specific antibody or antibody fragment which has been labeled with an appropriate detectable imaging moiety, such as a radioisotope (for example, ¹³¹I, ¹¹²In, ^{99m}Tc, (¹³¹I, ¹²⁵I, ¹²³I, ¹²¹I), carbon (¹⁴C), sulfur (³⁵S), tritium (³H), indium (^{115m}In, ^{113m}In, ¹¹²In, ¹¹¹In),
15 and technetium (⁹⁹Tc, ^{99m}Tc), thallium (²⁰¹Tl), gallium (⁶⁸Ga, ⁶⁷Ga), palladium (¹⁰³Pd), molybdenum (⁹⁹Mo), xenon (¹³³Xe), fluorine (¹⁸F), ¹⁵³Sm, ¹⁷⁷Lu, ¹⁵⁹Gd, ¹⁴⁹Pm, ¹⁴⁰La, ¹⁷⁵Yb, ¹⁶⁶Ho, ⁹⁰Y, ⁴⁷Sc, ¹⁸⁶Re, ¹⁸⁸Re, ¹⁴²Pr, ¹⁰⁵Rh, ⁹⁷Ru), a radio-opaque substance, or a material detectable by nuclear magnetic resonance, is introduced (for example, parenterally, subcutaneously or intraperitoneally) into the mammal to be examined for immune system
20 disorder. It will be understood in the art that the size of the subject and the imaging system used will determine the quantity of imaging moiety needed to produce diagnostic images. In the case of a radioisotope moiety, for a human subject, the quantity of radioactivity injected will normally range from about 5 to 20 millicuries of ^{99m}Tc. The labeled antibody or antibody fragment will then preferentially accumulate at the location of cells which express
25 the polypeptide encoded by a polynucleotide of the invention. *In vivo* tumor imaging is described in S.W. Burchiel et al., "Immunopharmacokinetics of Radiolabeled Antibodies and Their Fragments" (Chapter 13 in *Tumor Imaging: The Radiochemical Detection of Cancer*, S.W. Burchiel and B. A. Rhodes, eds., Masson Publishing Inc. (1982)).

30 In one embodiment, the invention provides a method for the specific delivery of compositions of the invention to cells by administering polypeptides of the invention (e.g., polypeptides encoded by polynucleotides of the invention and/or antibodies) that are associated with heterologous polypeptides or nucleic acids. In one example, the invention

provides a method for delivering a therapeutic protein into the targeted cell. In another example, the invention provides a method for delivering a single stranded nucleic acid (e.g., antisense or ribozymes) or double stranded nucleic acid (e.g., DNA that can integrate into the cell's genome or replicate episomally and that can be transcribed) into the targeted cell.

5 In another embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of tumor cells) by administering polypeptides of the invention in association with toxins or cytotoxic prodrugs.

By "toxin" is meant one or more compounds that bind and activate endogenous cytotoxic effector systems, radioisotopes, holotoxins, modified toxins, catalytic subunits of
10 toxins, or any molecules or enzymes not normally present in or on the surface of a cell that under defined conditions cause the cell's death. Toxins that may be used according to the methods of the invention include, but are not limited to, radioisotopes known in the art, compounds such as, for example, antibodies (or complement fixing containing portions thereof) that bind an inherent or induced endogenous cytotoxic effector system, thymidine
15 kinase, endonuclease, RNase, alpha toxin, ricin, abrin, *Pseudomonas* exotoxin A, diphtheria toxin, saporin, momordin, gelonin, pokeweed antiviral protein, alpha-sarcin and cholera toxin. "Toxin" also includes a cytostatic or cytocidal agent, a therapeutic agent or a radioactive metal ion, e.g., alpha-emitters such as, for example, ²¹³Bi, or other radioisotopes such as, for example, ¹⁰³Pd, ¹³³Xe, ¹³¹I, ⁶⁸Ge, ⁵⁷Co, ⁶⁵Zn, ⁸⁵Sr, ³²P, ³⁵S, ⁹⁰Y, ¹⁵³Sm, ¹⁵³Gd,
20 ¹⁶⁹Yb, ⁵¹Cr, ⁵⁴Mn, ⁷⁵Se, ¹¹³Sn, ⁹⁰Yttrium, ¹¹⁷Tin, ¹⁸⁶Rhenium, ¹⁶⁶Holmium, and ¹⁸⁸Rhenium; luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

Techniques known in the art may be applied to label polypeptides of the invention (including antibodies). Such techniques include, but are not limited to, the use of
25 bifunctional conjugating agents (see e.g., U.S. Patent Nos. 5,756,065; 5,714,631; 5,696,239; 5,652,361; 5,505,931; 5,489,425; 5,435,990; 5,428,139; 5,342,604; 5,274,119; 4,994,560; and 5,808,003; the contents of each of which are hereby incorporated by reference in its entirety).

Thus, the invention provides a diagnostic method of a disorder, which involves (a)
30 assaying the expression level of a polypeptide of the present invention in cells or body fluid of an individual; and (b) comparing the assayed polypeptide expression level with a standard polypeptide expression level, whereby an increase or decrease in the assayed polypeptide

expression level compared to the standard expression level is indicative of a disorder. With respect to cancer, the presence of a relatively high amount of transcript in biopsied tissue from an individual may indicate a predisposition for the development of the disease, or may provide a means for detecting the disease prior to the appearance of actual clinical symptoms.

5 A more definitive diagnosis of this type may allow health professionals to employ preventative measures or aggressive treatment earlier thereby preventing the development or further progression of the cancer.

Moreover, polypeptides of the present invention can be used to treat or prevent diseases or conditions such as, for example, neural disorders, immune system disorders,
10 muscular disorders, reproductive disorders, gastrointestinal disorders, pulmonary disorders, cardiovascular disorders, renal disorders, proliferative disorders, and/or cancerous diseases and conditions. For example, patients can be administered a polypeptide of the present invention in an effort to replace absent or decreased levels of the polypeptide (e.g., insulin), to supplement absent or decreased levels of a different polypeptide (e.g., hemoglobin S for
15 hemoglobin B, SOD, catalase, DNA repair proteins), to inhibit the activity of a polypeptide (e.g., an oncogene or tumor suppressor), to activate the activity of a polypeptide (e.g., by binding to a receptor), to reduce the activity of a membrane bound receptor by competing with it for free ligand (e.g., soluble TNF receptors used in reducing inflammation), or to bring about a desired response (e.g., blood vessel growth inhibition, enhancement of the
20 immune response to proliferative cells or tissues).

Similarly, antibodies directed to a polypeptide of the present invention can also be used to treat disease (as described supra, and elsewhere herein). For example, administration of an antibody directed to a polypeptide of the present invention can bind, and/or neutralize the polypeptide, and/or reduce overproduction of the polypeptide. Similarly, administration
25 of an antibody can activate the polypeptide, such as by binding to a polypeptide bound to a membrane (receptor).

At the very least, the polypeptides of the present invention can be used as molecular weight markers on SDS-PAGE gels or on molecular sieve gel filtration columns using methods well known to those of skill in the art. Polypeptides can also be used to raise
30 antibodies, which in turn are used to measure protein expression from a recombinant cell, as a way of assessing transformation of the host cell. Moreover, the polypeptides of the present invention can be used to test the following biological activities.

Gene Therapy Methods

Another aspect of the present invention is to gene therapy methods for treating or preventing disorders, diseases and conditions. The gene therapy methods relate to the introduction of nucleic acid (DNA, RNA and antisense DNA or RNA) sequences into an animal to achieve expression of the polypeptide of the present invention. This method requires a polynucleotide which codes for a polypeptide of the present invention operatively linked to a promoter and any other genetic elements necessary for the expression of the polypeptide by the target tissue. Such gene therapy and delivery techniques are known in the art, see, for example, WO90/11092, which is herein incorporated by reference.

Thus, for example, cells from a patient may be engineered with a polynucleotide (DNA or RNA) comprising a promoter operably linked to a polynucleotide of the present invention ex vivo, with the engineered cells then being provided to a patient to be treated with the polypeptide of the present invention. Such methods are well-known in the art. For example, see Beldegrun, A., et al., J. Natl. Cancer Inst. 85: 207-216 (1993); Ferrantini, M. et al., Cancer Research 53: 1107-1112 (1993); Ferrantini, M. et al., J. Immunology 153: 4604-4615 (1994); Kaido, T., et al., Int. J. Cancer 60: 221-229 (1995); Ogura, H., et al., Cancer Research 50: 5102-5106 (1990); Santodonato, L., et al., Human Gene Therapy 7:1-10 (1996); Santodonato, L., et al., Gene Therapy 4:1246-1255 (1997); and Zhang, J.-F. et al., Cancer Gene Therapy 3: 31-38 (1996)), which are herein incorporated by reference. In one embodiment, the cells which are engineered are arterial cells. The arterial cells may be reintroduced into the patient through direct injection to the artery, the tissues surrounding the artery, or through catheter injection.

As discussed in more detail below, the polynucleotide constructs can be delivered by any method that delivers injectable materials to the cells of an animal, such as, injection into the interstitial space of tissues (heart, muscle, skin, lung, liver, and the like). The polynucleotide constructs may be delivered in a pharmaceutically acceptable liquid or aqueous carrier.

In one embodiment, the polynucleotide of the present invention is delivered as a naked polynucleotide. The term "naked" polynucleotide, DNA or RNA refers to sequences that are free from any delivery vehicle that acts to assist, promote or facilitate entry into the cell, including viral sequences, viral particles, liposome formulations, lipofectin or

precipitating agents and the like. However, the polynucleotide of the present invention can also be delivered in liposome formulations and lipofectin formulations and the like can be prepared by methods well known to those skilled in the art. Such methods are described, for example, in U.S. Patent Nos. 5,593,972, 5,589,466, and 5,580,859, which are herein
5 incorporated by reference.

The polynucleotide vector constructs used in the gene therapy method are preferably constructs that will not integrate into the host genome nor will they contain sequences that allow for replication. Appropriate vectors include pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; pSVK3, pBPV, pMSG and pSVL available from Pharmacia;
10 and pEF1/V5, pcDNA3.1, and pRc/CMV2 available from Invitrogen. Other suitable vectors will be readily apparent to the skilled artisan.

Any strong promoter known to those skilled in the art can be used for driving the expression of the polynucleotide sequence. Suitable promoters include adenoviral promoters, such as the adenoviral major late promoter; or heterologous promoters, such as the
15 cytomegalovirus (CMV) promoter; the respiratory syncytial virus (RSV) promoter; inducible promoters, such as the MMT promoter, the metallothionein promoter; heat shock promoters; the albumin promoter; the ApoA1 promoter; human globin promoters; viral thymidine kinase promoters, such as the Herpes Simplex thymidine kinase promoter; retroviral LTRs; the β -actin promoter; and human growth hormone promoters. The promoter also may be the native
20 promoter for the polynucleotide of the present invention.

Unlike other gene therapy techniques, one major advantage of introducing naked nucleic acid sequences into target cells is the transitory nature of the polynucleotide synthesis in the cells. Studies have shown that non-replicating DNA sequences can be introduced into cells to provide production of the desired polypeptide for periods of up to six months.

The polynucleotide construct can be delivered to the interstitial space of tissues within
25 the an animal, including of muscle, skin, brain, lung, liver, spleen, bone marrow, thymus, heart, lymph, blood, bone, cartilage, pancreas, kidney, gall bladder, stomach, intestine, testis, ovary, uterus, rectum, nervous system, eye, gland, and connective tissue. Interstitial space of the tissues comprises the intercellular, fluid, mucopolysaccharide matrix among the reticular
30 fibers of organ tissues, elastic fibers in the walls of vessels or chambers, collagen fibers of fibrous tissues, or that same matrix within connective tissue ensheathing muscle cells or in the lacunae of bone. It is similarly the space occupied by the plasma of the circulation and the

lymph fluid of the lymphatic channels. Delivery to the interstitial space of muscle tissue is preferred for the reasons discussed below. They may be conveniently delivered by injection into the tissues comprising these cells. They are preferably delivered to and expressed in persistent, non-dividing cells which are differentiated, although delivery and expression may be achieved in non-differentiated or less completely differentiated cells, such as, for example, stem cells of blood or skin fibroblasts. In vivo muscle cells are particularly competent in their ability to take up and express polynucleotides.

For the naked nucleic acid sequence injection, an effective dosage amount of DNA or RNA will be in the range of from about 0.05 mg/kg body weight to about 50 mg/kg body weight. Preferably the dosage will be from about 0.005 mg/kg to about 20 mg/kg and more preferably from about 0.05 mg/kg to about 5 mg/kg. Of course, as the artisan of ordinary skill will appreciate, this dosage will vary according to the tissue site of injection. The appropriate and effective dosage of nucleic acid sequence can readily be determined by those of ordinary skill in the art and may depend on the condition being treated and the route of administration.

The preferred route of administration is by the parenteral route of injection into the interstitial space of tissues. However, other parenteral routes may also be used, such as, inhalation of an aerosol formulation particularly for delivery to lungs or bronchial tissues, throat or mucous membranes of the nose. In addition, naked DNA constructs can be delivered to arteries during angioplasty by the catheter used in the procedure.

The naked polynucleotides are delivered by any method known in the art, including, but not limited to, direct needle injection at the delivery site, intravenous injection, topical administration, catheter infusion, and so-called "gene guns". These delivery methods are known in the art.

The constructs may also be delivered with delivery vehicles such as viral sequences, viral particles, liposome formulations, lipofectin, precipitating agents, etc. Such methods of delivery are known in the art.

In certain embodiments, the polynucleotide constructs are complexed in a liposome preparation. Liposomal preparations for use in the instant invention include cationic (positively charged), anionic (negatively charged) and neutral preparations. However, cationic liposomes are particularly preferred because a tight charge complex can be formed between the cationic liposome and the polyanionic nucleic acid. Cationic liposomes have

been shown to mediate intracellular delivery of plasmid DNA (Felgner et al., Proc. Natl. Acad. Sci. USA (1987) 84:7413-7416, which is herein incorporated by reference); mRNA (Malone et al., Proc. Natl. Acad. Sci. USA (1989) 86:6077-6081, which is herein incorporated by reference); and purified transcription factors (Debs et al., J. Biol. Chem. (1990) 265:10189-10192, which is herein incorporated by reference), in functional form.

Cationic liposomes are readily available. For example, N[1-2,3-dioleoyloxy)propyl]-N,N,N-triethylammonium (DOTMA) liposomes are particularly useful and are available under the trademark Lipofectin, from GIBCO BRL, Grand Island, N.Y. (See, also, Felgner et al., Proc. Natl Acad. Sci. USA (1987) 84:7413-7416, which is herein incorporated by reference). Other commercially available liposomes include transfectace (DDAB/DOPE) and DOTAP/DOPE (Boehringer).

Other cationic liposomes can be prepared from readily available materials using techniques well known in the art. See, e.g. PCT Publication No. WO 90/11092 (which is herein incorporated by reference) for a description of the synthesis of DOTAP (1,2-bis(oleoyloxy)-3-(trimethylammonio)propane) liposomes. Preparation of DOTMA liposomes is explained in the literature, see, e.g., P. Felgner et al., Proc. Natl. Acad. Sci. USA 84:7413-7417, which is herein incorporated by reference. Similar methods can be used to prepare liposomes from other cationic lipid materials.

Similarly, anionic and neutral liposomes are readily available, such as from Avanti Polar Lipids (Birmingham, Ala.), or can be easily prepared using readily available materials. Such materials include phosphatidyl, choline, cholesterol, phosphatidyl ethanolamine, dioleoylphosphatidyl choline (DOPC), dioleoylphosphatidyl glycerol (DOPG), dioleoylphosphatidyl ethanolamine (DOPE), among others. These materials can also be mixed with the DOTMA and DOTAP starting materials in appropriate ratios. Methods for making liposomes using these materials are well known in the art.

For example, commercially dioleoylphosphatidyl choline (DOPC), dioleoylphosphatidyl glycerol (DOPG), and dioleoylphosphatidyl ethanolamine (DOPE) can be used in various combinations to make conventional liposomes, with or without the addition of cholesterol. Thus, for example, DOPG/DOPC vesicles can be prepared by drying 50 mg each of DOPG and DOPC under a stream of nitrogen gas into a sonication vial. The sample is placed under a vacuum pump overnight and is hydrated the following day with deionized water. The sample is then sonicated for 2 hours in a capped vial, using a Heat

Systems model 350 sonicator equipped with an inverted cup (bath type) probe at the maximum setting while the bath is circulated at 15EC. Alternatively, negatively charged vesicles can be prepared without sonication to produce multilamellar vesicles or by extrusion through nucleopore membranes to produce unilamellar vesicles of discrete size. Other methods are known and available to those of skill in the art.

The liposomes can comprise multilamellar vesicles (MLVs), small unilamellar vesicles (SUVs), or large unilamellar vesicles (LUVs), with SUVs being preferred. The various liposome-nucleic acid complexes are prepared using methods well known in the art. See, e.g., Straubinger et al., *Methods of Immunology* (1983), 101:512-527, which is herein incorporated by reference. For example, MLVs containing nucleic acid can be prepared by depositing a thin film of phospholipid on the walls of a glass tube and subsequently hydrating with a solution of the material to be encapsulated. SUVs are prepared by extended sonication of MLVs to produce a homogeneous population of unilamellar liposomes. The material to be entrapped is added to a suspension of preformed MLVs and then sonicated. When using liposomes containing cationic lipids, the dried lipid film is resuspended in an appropriate solution such as sterile water or an isotonic buffer solution such as 10 mM Tris/NaCl, sonicated, and then the preformed liposomes are mixed directly with the DNA. The liposome and DNA form a very stable complex due to binding of the positively charged liposomes to the cationic DNA. SUVs find use with small nucleic acid fragments. LUVs are prepared by a number of methods, well known in the art. Commonly used methods include Ca^{2+} -EDTA chelation (Papahadjopoulos et al., *Biochim. Biophys. Acta* (1975) 394:483; Wilson et al., *Cell* (1979) 17:77); ether injection (Deamer, D. and Bangham, A., *Biochim. Biophys. Acta* (1976) 443:629; Ostro et al., *Biochem. Biophys. Res. Commun.* (1977) 76:836; Fraley et al., *Proc. Natl. Acad. Sci. USA* (1979) 76:3348); detergent dialysis (Enoch, H. and Strittmatter, P., *Proc. Natl. Acad. Sci. USA* (1979) 76:145); and reverse-phase evaporation (REV) (Fraley et al., *J. Biol. Chem.* (1980) 255:10431; Szoka, F. and Papahadjopoulos, D., *Proc. Natl. Acad. Sci. USA* (1978) 75:145; Schaefer-Ridder et al., *Science* (1982) 215:166), which are herein incorporated by reference.

Generally, the ratio of DNA to liposomes will be from about 10:1 to about 1:10. Preferably, the ration will be from about 5:1 to about 1:5. More preferably, the ration will be about 3:1 to about 1:3. Still more preferably, the ratio will be about 1:1.

U.S. Patent No. 5,676,954 (which is herein incorporated by reference) reports on the injection of genetic material, complexed with cationic liposomes carriers, into mice. U.S. Patent Nos. 4,897,355, 4,946,787, 5,049,386, 5,459,127, 5,589,466, 5,693,622, 5,580,859, 5,703,055, and international publication no. WO 94/9469 (which are herein incorporated by reference) provide cationic lipids for use in transfecting DNA into cells and mammals. U.S. Patent Nos. 5,589,466, 5,693,622, 5,580,859, 5,703,055, and international publication no. WO 94/9469 (which are herein incorporated by reference) provide methods for delivering DNA-cationic lipid complexes to mammals.

In certain embodiments, cells are engineered, ex vivo or in vivo, using a retroviral particle containing RNA which comprises a sequence encoding a polypeptide of the present invention. Retroviruses from which the retroviral plasmid vectors may be derived include, but are not limited to, Moloney Murine Leukemia Virus, spleen necrosis virus, Rous sarcoma Virus, Harvey Sarcoma Virus, avian leukosis virus, gibbon ape leukemia virus, human immunodeficiency virus, Myeloproliferative Sarcoma Virus, and mammary tumor virus.

The retroviral plasmid vector is employed to transduce packaging cell lines to form producer cell lines. Examples of packaging cells which may be transfected include, but are not limited to, the PE501, PA317, R-2, R-AM, PA12, T19-14X, VT-19-17-H2, RCRE, RCRIP, GP+E-86, GP+envAm12, and DAN cell lines as described in Miller, Human Gene Therapy 1:5-14 (1990), which is incorporated herein by reference in its entirety. The vector may transduce the packaging cells through any means known in the art. Such means include, but are not limited to, electroporation, the use of liposomes, and CaPO_4 precipitation. In one alternative, the retroviral plasmid vector may be encapsulated into a liposome, or coupled to a lipid, and then administered to a host.

The producer cell line generates infectious retroviral vector particles which include polynucleotide encoding a polypeptide of the present invention. Such retroviral vector particles then may be employed, to transduce eukaryotic cells, either in vitro or in vivo. The transduced eukaryotic cells will express a polypeptide of the present invention.

In certain other embodiments, cells are engineered, ex vivo or in vivo, with polynucleotide contained in an adenovirus vector. Adenovirus can be manipulated such that it encodes and expresses a polypeptide of the present invention, and at the same time is inactivated in terms of its ability to replicate in a normal lytic viral life cycle. Adenovirus expression is achieved without integration of the viral DNA into the host cell chromosome,

thereby alleviating concerns about insertional mutagenesis. Furthermore, adenoviruses have been used as live enteric vaccines for many years with an excellent safety profile (Schwartz, A. R. et al. (1974) *Am. Rev. Respir. Dis.* 109:233-238). Finally, adenovirus mediated gene transfer has been demonstrated in a number of instances including transfer of
5 alpha-1-antitrypsin and CFTR to the lungs of cotton rats (Rosenfeld, M. A. et al. (1991) *Science* 252:431-434; Rosenfeld et al., (1992) *Cell* 68:143-155). Furthermore, extensive studies to attempt to establish adenovirus as a causative agent in human cancer were uniformly negative (Green, M. et al. (1979) *Proc. Natl. Acad. Sci. USA* 76:6606).

Suitable adenoviral vectors useful in the present invention are described, for example,
10 in Kozarsky and Wilson, *Curr. Opin. Genet. Devel.* 3:499-503 (1993); Rosenfeld et al., *Cell* 68:143-155 (1992); Engelhardt et al., *Human Genet. Ther.* 4:759-769 (1993); Yang et al., *Nature Genet.* 7:362-369 (1994); Wilson et al., *Nature* 365:691-692 (1993); and U.S. Patent No. 5,652,224, which are herein incorporated by reference. For example, the adenovirus vector Ad2 is useful and can be grown in human 293 cells. These cells contain the E1 region
15 of adenovirus and constitutively express E1a and E1b, which complement the defective adenoviruses by providing the products of the genes deleted from the vector. In addition to Ad2, other varieties of adenovirus (e.g., Ad3, Ad5, and Ad7) are also useful in the present invention.

Preferably, the adenoviruses used in the present invention are replication deficient.
20 Replication deficient adenoviruses require the aid of a helper virus and/or packaging cell line to form infectious particles. The resulting virus is capable of infecting cells and can express a polynucleotide of interest which is operably linked to a promoter, but cannot replicate in most cells. Replication deficient adenoviruses may be deleted in one or more of all or a portion of the following genes: E1a, E1b, E3, E4, E2a, or L1 through L5.

25 In certain other embodiments, the cells are engineered, ex vivo or in vivo, using an adeno-associated virus (AAV). AAVs are naturally occurring defective viruses that require helper viruses to produce infectious particles (Muzyczka, N., *Curr. Topics in Microbiol. Immunol.* 158:97 (1992)). It is also one of the few viruses that may integrate its DNA into non-dividing cells. Vectors containing as little as 300 base pairs of AAV can be packaged and
30 can integrate, but space for exogenous DNA is limited to about 4.5 kb. Methods for producing and using such AAVs are known in the art. See, for example, U.S. Patent Nos. 5,139,941, 5,173,414, 5,354,678, 5,436,146, 5,474,935, 5,478,745, and 5,589,377.

For example, an appropriate AAV vector for use in the present invention will include all the sequences necessary for DNA replication, encapsidation, and host-cell integration. The polynucleotide construct is inserted into the AAV vector using standard cloning methods, such as those found in Sambrook et al., *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Press (1989). The recombinant AAV vector is then transfected into packaging cells which are infected with a helper virus, using any standard technique, including lipofection, electroporation, calcium phosphate precipitation, etc. Appropriate helper viruses include adenoviruses, cytomegaloviruses, vaccinia viruses, or herpes viruses. Once the packaging cells are transfected and infected, they will produce infectious AAV viral particles which contain the polynucleotide construct. These viral particles are then used to transduce eukaryotic cells, either ex vivo or in vivo. The transduced cells will contain the polynucleotide construct integrated into its genome, and will express a polypeptide of the invention.

Another method of gene therapy involves operably associating heterologous control regions and endogenous polynucleotide sequences (e.g. encoding a polypeptide of the present invention) via homologous recombination (see, e.g., U.S. Patent No. 5,641,670, issued June 24, 1997; International Publication No. WO 96/29411, published September 26, 1996; International Publication No. WO 94/12650, published August 4, 1994; Koller et al., *Proc. Natl. Acad. Sci. USA* 86:8932-8935 (1989); and Zijlstra et al., *Nature* 342:435-438 (1989). This method involves the activation of a gene which is present in the target cells, but which is not normally expressed in the cells, or is expressed at a lower level than desired.

Polynucleotide constructs are made, using standard techniques known in the art, which contain the promoter with targeting sequences flanking the promoter. Suitable promoters are described herein. The targeting sequence is sufficiently complementary to an endogenous sequence to permit homologous recombination of the promoter-targeting sequence with the endogenous sequence. The targeting sequence will be sufficiently near the 5' end of the desired endogenous polynucleotide sequence so the promoter will be operably linked to the endogenous sequence upon homologous recombination.

The promoter and the targeting sequences can be amplified using PCR. Preferably, the amplified promoter contains distinct restriction enzyme sites on the 5' and 3' ends. Preferably, the 3' end of the first targeting sequence contains the same restriction enzyme site as the 5' end of the amplified promoter and the 5' end of the second targeting sequence

contains the same restriction site as the 3' end of the amplified promoter. The amplified promoter and targeting sequences are digested and ligated together.

The promoter-targeting sequence construct is delivered to the cells, either as naked polynucleotide, or in conjunction with transfection-facilitating agents, such as liposomes, viral sequences, viral particles, whole viruses, lipofection, precipitating agents, etc., described in more detail above. The P promoter-targeting sequence can be delivered by any method, included direct needle injection, intravenous injection, topical administration, catheter infusion, particle accelerators, etc. The methods are described in more detail below.

The promoter-targeting sequence construct is taken up by cells. Homologous recombination between the construct and the endogenous sequence takes place, such that an endogenous sequence is placed under the control of the promoter. The promoter then drives the expression of the endogenous sequence.

The polynucleotides encoding a polypeptide of the present invention may be administered along with other polynucleotides encoding an angiogenic protein. Examples of angiogenic proteins include, but are not limited to, acidic and basic fibroblast growth factors, VEGF-1, VEGF-2, VEGF-3, epidermal growth factor alpha and beta, platelet-derived endothelial cell growth factor, platelet-derived growth factor, tumor necrosis factor alpha, hepatocyte growth factor, insulin like growth factor, colony stimulating factor, macrophage colony stimulating factor, granulocyte/macrophage colony stimulating factor, and nitric oxide synthase.

Preferably, the polynucleotide encoding a polypeptide of the present invention contains a secretory signal sequence that facilitates secretion of the protein. Typically, the signal sequence is positioned in the coding region of the polynucleotide to be expressed towards or at the 5' end of the coding region. The signal sequence may be homologous or heterologous to the polynucleotide of interest and may be homologous or heterologous to the cells to be transfected. Additionally, the signal sequence may be chemically synthesized using methods known in the art.

Any mode of administration of any of the above-described polynucleotides constructs can be used so long as the mode results in the expression of one or more molecules in an amount sufficient to provide a therapeutic effect. This includes direct needle injection, systemic injection, catheter infusion, biolistic injectors, particle accelerators (i.e., "gene guns"), gelfoam sponge depots, other commercially available depot materials, osmotic pumps

(e.g., Alza minipumps), oral or suppository solid (tablet or pill) pharmaceutical formulations, and decanting or topical applications during surgery. For example, direct injection of naked calcium phosphate-precipitated plasmid into rat liver and rat spleen or a protein-coated plasmid into the portal vein has resulted in gene expression of the foreign gene in the rat livers (Kaneda et al., Science 243:375 (1989)).

A preferred method of local administration is by direct injection. Preferably, a recombinant molecule of the present invention complexed with a delivery vehicle is administered by direct injection into or locally within the area of arteries. Administration of a composition locally within the area of arteries refers to injecting the composition centimeters and preferably, millimeters within arteries.

Another method of local administration is to contact a polynucleotide construct of the present invention in or around a surgical wound. For example, a patient can undergo surgery and the polynucleotide construct can be coated on the surface of tissue inside the wound or the construct can be injected into areas of tissue inside the wound.

Therapeutic compositions useful in systemic administration, include recombinant molecules of the present invention complexed to a targeted delivery vehicle of the present invention. Suitable delivery vehicles for use with systemic administration comprise liposomes comprising ligands for targeting the vehicle to a particular site.

Preferred methods of systemic administration, include intravenous injection, aerosol, oral and percutaneous (topical) delivery. Intravenous injections can be performed using methods standard in the art. Aerosol delivery can also be performed using methods standard in the art (see, for example, Stribling et al., Proc. Natl. Acad. Sci. USA 189:11277-11281, 1992, which is incorporated herein by reference). Oral delivery can be performed by complexing a polynucleotide construct of the present invention to a carrier capable of withstanding degradation by digestive enzymes in the gut of an animal. Examples of such carriers, include plastic capsules or tablets, such as those known in the art. Topical delivery can be performed by mixing a polynucleotide construct of the present invention with a lipophilic reagent (e.g., DMSO) that is capable of passing into the skin.

Determining an effective amount of substance to be delivered can depend upon a number of factors including, for example, the chemical structure and biological activity of the substance, the age and weight of the animal, the precise condition requiring treatment and its severity, and the route of administration. The frequency of treatments depends upon a

number of factors, such as the amount of polynucleotide constructs administered per dose, as well as the health and history of the subject. The precise amount, number of doses, and timing of doses will be determined by the attending physician or veterinarian.

Therapeutic compositions of the present invention can be administered to any animal, preferably to mammals and birds. Preferred mammals include humans, dogs, cats, mice, rats, rabbits sheep, cattle, horses and pigs, with humans being particularly preferred.

Biological Activities

Polynucleotides or polypeptides, or agonists or antagonists of the present invention, can be used in assays to test for one or more biological activities. If these polynucleotides or polypeptides, or agonists or antagonists of the present invention, do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides and polypeptides, and agonists or antagonists could be used to treat the associated disease.

Immune Activity

A polypeptide or polynucleotide, or agonists or antagonists of the present invention may be useful in treating deficiencies or disorders of the immune system, by activating or inhibiting the proliferation, differentiation, or mobilization (chemotaxis) of immune cells. Immune cells develop through a process called hematopoiesis, producing myeloid (platelets, red blood cells, neutrophils, and macrophages) and lymphoid (B and T lymphocytes) cells from pluripotent stem cells. The etiology of these immune deficiencies or disorders may be genetic, somatic, such as cancer or some autoimmune disorders, acquired (e.g., by chemotherapy or toxins), or infectious. Moreover, polynucleotides or polypeptides, or agonists or antagonists of the present invention can be used as a marker or detector of a particular immune system disease or disorder.

Polynucleotides or polypeptides, or agonists or antagonists of the present invention may be useful in treating or detecting deficiencies or disorders of hematopoietic cells. Polynucleotides or polypeptides, or agonists or antagonists of the present invention could be used to increase differentiation and proliferation of hematopoietic cells, including the pluripotent stem cells, in an effort to treat those disorders associated with a decrease in certain (or many) types hematopoietic cells. Examples of immunologic deficiency

syndromes include, but are not limited to: blood protein disorders (e.g. agammaglobulinemia, dysgammaglobulinemia), ataxia telangiectasia, common variable immunodeficiency, Digeorge Syndrome, HIV infection, HTLV-BLV infection, leukocyte adhesion deficiency syndrome, lymphopenia, phagocyte bactericidal dysfunction, severe
5 combined immunodeficiency (SCIDs), Wiskott-Aldrich Disorder, anemia, thrombocytopenia, or hemoglobinuria.

Moreover, polynucleotides or polypeptides, or agonists or antagonists of the present invention could also be used to modulate hemostatic (the stopping of bleeding) or thrombolytic activity (clot formation). For example, by increasing hemostatic or
10 thrombolytic activity, polynucleotides or polypeptides, or agonists or antagonists of the present invention could be used to treat blood coagulation disorders (e.g., afibrinogenemia, factor deficiencies), blood platelet disorders (e.g. thrombocytopenia), or wounds resulting from trauma, surgery, or other causes. Alternatively, polynucleotides or polypeptides, or agonists or antagonists of the present invention that can decrease hemostatic or thrombolytic
15 activity could be used to inhibit or dissolve clotting. These molecules could be important in the treatment of heart attacks (infarction), strokes, or scarring.

Polynucleotides or polypeptides, or agonists or antagonists of the present invention may also be useful in treating or detecting autoimmune disorders. Many autoimmune disorders result from inappropriate recognition of self as foreign material by immune cells.
20 This inappropriate recognition results in an immune response leading to the destruction of the host tissue. Therefore, the administration of polynucleotides or polypeptides, or agonists or antagonists of the present invention that can inhibit an immune response, particularly the proliferation, differentiation, or chemotaxis of T-cells, may be an effective therapy in preventing autoimmune disorders.

25 Examples of autoimmune disorders that can be treated or detected include, but are not limited to: Addison's Disease, hemolytic anemia, antiphospholipid syndrome, rheumatoid arthritis, dermatitis, allergic encephalomyelitis, glomerulonephritis, Goodpasture's Syndrome, Graves' Disease, Multiple Sclerosis, Myasthenia Gravis, Neuritis, Ophthalmia, Bullous Pemphigoid, Pemphigus, Polyendocrinopathies, Purpura, Reiter's Disease, Stiff-Man
30 Syndrome, Autoimmune Thyroiditis, Systemic Lupus Erythematosus, Autoimmune Pulmonary Inflammation, Guillain-Barre Syndrome, insulin dependent diabetes mellitus, and autoimmune inflammatory eye disease.

Similarly, allergic reactions and conditions, such as asthma (particularly allergic asthma) or other respiratory problems, may also be treated by polynucleotides or polypeptides, or agonists or antagonists of the present invention. Moreover, these molecules can be used to treat anaphylaxis, hypersensitivity to an antigenic molecule, or blood group incompatibility.

Polynucleotides or polypeptides, or agonists or antagonists of the present invention may also be used to treat and/or prevent organ rejection or graft-versus-host disease (GVHD). Organ rejection occurs by host immune cell destruction of the transplanted tissue through an immune response. Similarly, an immune response is also involved in GVHD, but, in this case, the foreign transplanted immune cells destroy the host tissues. The administration of polynucleotides or polypeptides, or agonists or antagonists of the present invention that inhibits an immune response, particularly the proliferation, differentiation, or chemotaxis of T-cells, may be an effective therapy in preventing organ rejection or GVHD.

Similarly, polynucleotides or polypeptides, or agonists or antagonists of the present invention may also be used to modulate inflammation. For example, polynucleotides or polypeptides, or agonists or antagonists of the present invention may inhibit the proliferation and differentiation of cells involved in an inflammatory response. These molecules can be used to treat inflammatory conditions, both chronic and acute conditions, including chronic prostatitis, granulomatous prostatitis and malacoplakia, inflammation associated with infection (e.g., septic shock, sepsis, or systemic inflammatory response syndrome (SIRS)), ischemia-reperfusion injury, endotoxin lethality, arthritis, complement-mediated hyperacute rejection, nephritis, cytokine or chemokine induced lung injury, inflammatory bowel disease, Crohn's disease, or resulting from over production of cytokines (e.g., TNF or IL-1.)

Hyperproliferative Disorders

Polynucleotides or polypeptides, or agonists or antagonists of the present invention can be used to treat or detect hyperproliferative disorders, including neoplasms. Polynucleotides or polypeptides, or agonists or antagonists of the present invention may inhibit the proliferation of the disorder through direct or indirect interactions. Alternatively, Polynucleotides or polypeptides, or agonists or antagonists of the present invention may proliferate other cells which can inhibit the hyperproliferative disorder.

For example, by increasing an immune response, particularly increasing antigenic qualities of the hyperproliferative disorder or by proliferating, differentiating, or mobilizing T-cells, hyperproliferative disorders can be treated. This immune response may be increased by either enhancing an existing immune response, or by initiating a new immune response.

5 Alternatively, decreasing an immune response may also be a method of treating hyperproliferative disorders, such as a chemotherapeutic agent.

Examples of hyperproliferative disorders that can be treated or detected by Polynucleotides or polypeptides, or agonists or antagonists of the present invention include, but are not limited to neoplasms located in the: colon, abdomen, bone, breast, digestive

10 system, liver, pancreas, peritoneum, endocrine glands (adrenal, parathyroid, pituitary, testicles, ovary, thymus, thyroid), eye, head and neck, nervous (central and peripheral), lymphatic system, pelvic, skin, soft tissue, spleen, thoracic, and urogenital.

Similarly, other hyperproliferative disorders can also be treated or detected by polynucleotides or polypeptides, or agonists or antagonists of the present invention.

15 Examples of such hyperproliferative disorders include, but are not limited to: hypergammaglobulinemia, lymphoproliferative disorders, paraproteinemias, purpura, sarcoidosis, Sezary Syndrome, Waldenström's Macroglobulinemia, Gaucher's Disease, histiocytosis, and any other hyperproliferative disease, besides neoplasia, located in an organ system listed above.

20 One preferred embodiment utilizes polynucleotides of the present invention to inhibit aberrant cellular division, by gene therapy using the present invention, and/or protein fusions or fragments thereof.

Thus, the present invention provides a method for treating cell proliferative disorders by inserting into an abnormally proliferating cell a polynucleotide of the present invention,

25 wherein said polynucleotide represses said expression.

Another embodiment of the present invention provides a method of treating cell-proliferative disorders in individuals comprising administration of one or more active gene copies of the present invention to an abnormally proliferating cell or cells. In a preferred embodiment, polynucleotides of the present invention is a DNA construct comprising a

30 recombinant expression vector effective in expressing a DNA sequence encoding said polynucleotides. In another preferred embodiment of the present invention, the DNA construct encoding the polynucleotides of the present invention is inserted into cells to be

treated utilizing a retrovirus, or more preferably an adenoviral vector (See G J. Nabel, et. al., PNAS 1999 96: 324-326, which is hereby incorporated by reference). In a most preferred embodiment, the viral vector is defective and will not transform non-proliferating cells, only proliferating cells. Moreover, in a preferred embodiment, the polynucleotides of the present invention inserted into proliferating cells either alone, or in combination with or fused to other polynucleotides, can then be modulated via an external stimulus (i.e. magnetic, specific small molecule, chemical, or drug administration, etc.), which acts upon the promoter upstream of said polynucleotides to induce expression of the encoded protein product. As such the beneficial therapeutic affect of the present invention may be expressly modulated (i.e. to increase, decrease, or inhibit expression of the present invention) based upon said external stimulus.

Polynucleotides of the present invention may be useful in repressing expression of oncogenic genes or antigens. By "repressing expression of the oncogenic genes " is intended the suppression of the transcription of the gene, the degradation of the gene transcript (pre-message RNA), the inhibition of splicing, the destruction of the messenger RNA, the prevention of the post-translational modifications of the protein, the destruction of the protein, or the inhibition of the normal function of the protein.

For local administration to abnormally proliferating cells, polynucleotides of the present invention may be administered by any method known to those of skill in the art including, but not limited to transfection, electroporation, microinjection of cells, or in vehicles such as liposomes, lipofectin, or as naked polynucleotides, or any other method described throughout the specification. The polynucleotide of the present invention may be delivered by known gene delivery systems such as, but not limited to, retroviral vectors (Gilboa, J. Virology 44:845 (1982); Hocke, Nature 320:275 (1986); Wilson, et al., Proc. Natl. Acad. Sci. U.S.A. 85:3014), vaccinia virus system (Chakrabarty et al., Mol. Cell Biol. 5:3403 (1985) or other efficient DNA delivery systems (Yates et al., Nature 313:812 (1985)) known to those skilled in the art. These references are exemplary only and are hereby incorporated by reference. In order to specifically deliver or transfect cells which are abnormally proliferating and spare non-dividing cells, it is preferable to utilize a retrovirus, or adenoviral (as described in the art and elsewhere herein) delivery system known to those of skill in the art. Since host DNA replication is required for retroviral DNA to integrate and the retrovirus will be unable to self replicate due to the lack of the retrovirus genes needed for its life cycle.

Utilizing such a retroviral delivery system for polynucleotides of the present invention will target said gene and constructs to abnormally proliferating cells and will spare the non-dividing normal cells.

5 The polynucleotides of the present invention may be delivered directly to cell proliferative disorder/disease sites in internal organs, body cavities and the like by use of imaging devices used to guide an injecting needle directly to the disease site. The polynucleotides of the present invention may also be administered to disease sites at the time of surgical intervention.

10 By "cell proliferative disease" is meant any human or animal disease or disorder, affecting any one or any combination of organs, cavities, or body parts, which is characterized by single or multiple local abnormal proliferations of cells, groups of cells, or tissues, whether benign or malignant.

Any amount of the polynucleotides of the present invention may be administered as long as it has a biologically inhibiting effect on the proliferation of the treated cells. 15 Moreover, it is possible to administer more than one of the polynucleotide of the present invention simultaneously to the same site. By "biologically inhibiting" is meant partial or total growth inhibition as well as decreases in the rate of proliferation or growth of the cells. The biologically inhibitory dose may be determined by assessing the effects of the polynucleotides of the present invention on target malignant or abnormally proliferating cell 20 growth in tissue culture, tumor growth in animals and cell cultures, or any other method known to one of ordinary skill in the art.

The present invention is further directed to antibody-based therapies which involve administering of anti-polypeptides and anti-polynucleotide antibodies to a mammalian, preferably human, patient for treating one or more of the described disorders. Methods for 25 producing anti-polypeptides and anti-polynucleotide antibodies polyclonal and monoclonal antibodies are described in detail elsewhere herein. Such antibodies may be provided in pharmaceutically acceptable compositions as known in the art or as described herein.

A summary of the ways in which the antibodies of the present invention may be used therapeutically includes binding polynucleotides or polypeptides of the present invention 30 locally or systemically in the body or by direct cytotoxicity of the antibody, e.g. as mediated by complement (CDC) or by effector cells (ADCC). Some of these approaches are described in more detail below. Armed with the teachings provided herein, one of ordinary skill in the

art will know how to use the antibodies of the present invention for diagnostic, monitoring or therapeutic purposes without undue experimentation.

In particular, the antibodies, fragments and derivatives of the present invention are useful for treating a subject having or developing cell proliferative and/or differentiation disorders as described herein. Such treatment comprises administering a single or multiple
5 doses of the antibody, or a fragment, derivative, or a conjugate thereof.

The antibodies of this invention may be advantageously utilized in combination with other monoclonal or chimeric antibodies, or with lymphokines or hematopoietic growth factors, for example., which serve to increase the number or activity of effector cells which
10 interact with the antibodies.

It is preferred to use high affinity and/or potent in vivo inhibiting and/or neutralizing antibodies against polypeptides or polynucleotides of the present invention, fragments or regions thereof, for both immunoassays directed to and therapy of disorders related to polynucleotides or polypeptides, including fragments thereof, of the present invention. Such
15 antibodies, fragments, or regions, will preferably have an affinity for polynucleotides or polypeptides, including fragments thereof. Preferred binding affinities include those with a dissociation constant or K_d less than $5 \times 10^{-6}M$, $10^{-6}M$, $5 \times 10^{-7}M$, $10^{-7}M$, $5 \times 10^{-8}M$, $10^{-8}M$, $5 \times 10^{-9}M$, $10^{-9}M$, $5 \times 10^{-10}M$, $10^{-10}M$, $5 \times 10^{-11}M$, $10^{-11}M$, $5 \times 10^{-12}M$, $10^{-12}M$, $5 \times 10^{-13}M$, $10^{-13}M$, $5 \times 10^{-14}M$, $10^{-14}M$, $5 \times 10^{-15}M$, and $10^{-15}M$.

Moreover, polypeptides of the present invention are useful in inhibiting the angiogenesis of proliferative cells or tissues, either alone, as a protein fusion, or in combination with other polypeptides directly or indirectly, as described elsewhere herein. In a most preferred embodiment, said anti-angiogenesis effect may be achieved indirectly, for example, through the inhibition of hematopoietic, tumor-specific cells, such as tumor-associated macrophages (See Joseph IB, et al. J Natl Cancer Inst, 90(21):1648-53 (1998),
25 which is hereby incorporated by reference). Antibodies directed to polypeptides or polynucleotides of the present invention may also result in inhibition of angiogenesis directly, or indirectly (See Witte L, et al., Cancer Metastasis Rev. 17(2):155-61 (1998), which is hereby incorporated by reference)).

Polypeptides, including protein fusions, of the present invention, or fragments thereof may be useful in inhibiting proliferative cells or tissues through the induction of apoptosis. Said polypeptides may act either directly, or indirectly to induce apoptosis of proliferative
30

cells and tissues, for example in the activation of a death-domain receptor, such as tumor necrosis factor (TNF) receptor-1, CD95 (Fas/APO-1), TNF-receptor-related apoptosis-mediated protein (TRAMP) and TNF-related apoptosis-inducing ligand (TRAIL) receptor-1 and -2 (See Schulze-Osthoff K, et.al., Eur J Biochem 254(3):439-59 (1998), which is hereby
5 incorporated by reference). Moreover, in another preferred embodiment of the present invention, said polypeptides may induce apoptosis through other mechanisms, such as in the activation of other proteins which will activate apoptosis, or through stimulating the expression of said proteins, either alone or in combination with small molecule drugs or adjuvants, such as apoptonin, galectins, thioredoxins, antiinflammatory proteins (See for
10 example, Mutat Res 400(1-2):447-55 (1998), Med Hypotheses.50(5):423-33 (1998), Chem Biol Interact. Apr 24;111-112:23-34 (1998), J Mol Med.76(6):402-12 (1998), Int J Tissue React;20(1):3-15 (1998), which are all hereby incorporated by reference).

Polypeptides, including protein fusions to, or fragments thereof, of the present invention are useful in inhibiting the metastasis of proliferative cells or tissues. Inhibition
15 may occur as a direct result of administering polypeptides, or antibodies directed to said polypeptides as described elsewhere herein, or indirectly, such as activating the expression of proteins known to inhibit metastasis, for example alpha 4 integrins, (See, e.g., Curr Top Microbiol Immunol 1998;231:125-41, which is hereby incorporated by reference). Such therapeutic affects of the present invention may be achieved either alone, or in combination
20 with small molecule drugs or adjuvants.

In another embodiment, the invention provides a method of delivering compositions containing the polypeptides of the invention (e.g., compositions containing polypeptides or polypeptide antibodies associated with heterologous polypeptides, heterologous nucleic acids, toxins, or prodrugs) to targeted cells expressing the polypeptide of the present invention.
25 Polypeptides or polypeptide antibodies of the invention may be associated with with heterologous polypeptides, heterologous nucleic acids, toxins, or prodrugs via hydrophobic, hydrophilic, ionic and/or covalent interactions.

Polypeptides, protein fusions to, or fragments thereof, of the present invention are useful in enhancing the immunogenicity and/or antigenicity of proliferating cells or tissues,
30 either directly, such as would occur if the polypeptides of the present invention 'vaccinated' the immune response to respond to proliferative antigens and immunogens, or indirectly, such

as in activating the expression of proteins known to enhance the immune response (e.g. chemokines), to said antigens and immunogens.

Cardiovascular Disorders

5 Polynucleotides or polypeptides, or agonists or antagonists of the present invention, may be used to treat cardiovascular disorders, including peripheral artery disease, such as limb ischemia.

Cardiovascular disorders include cardiovascular abnormalities, such as arterio-arterial fistula, arteriovenous fistula, cerebral arteriovenous malformations, congenital heart defects, 10 pulmonary atresia, and Scimitar Syndrome. Congenital heart defects include aortic coarctation, cor triatriatum, coronary vessel anomalies, crisscross heart, dextrocardia, patent ductus arteriosus, Ebstein's anomaly, Eisenmenger complex, hypoplastic left heart syndrome, levocardia, tetralogy of fallot, transposition of great vessels, double outlet right ventricle, tricuspid atresia, persistent truncus arteriosus, and heart septal defects, such as 15 aortopulmonary septal defect, endocardial cushion defects, Lutembacher's Syndrome, trilogy of Fallot, ventricular heart septal defects.

Cardiovascular disorders also include heart disease, such as arrhythmias, carcinoid heart disease, high cardiac output, low cardiac output, cardiac tamponade, endocarditis (including bacterial), heart aneurysm, cardiac arrest, congestive heart failure, congestive 20 cardiomyopathy, paroxysmal dyspnea, cardiac edema, heart hypertrophy, congestive cardiomyopathy, left ventricular hypertrophy, right ventricular hypertrophy, post-infarction heart rupture, ventricular septal rupture, heart valve diseases, myocardial diseases, myocardial ischemia, pericardial effusion, pericarditis (including constrictive and tuberculous), pneumopericardium, postpericardiectomy syndrome, pulmonary heart disease, 25 rheumatic heart disease, ventricular dysfunction, hyperemia, cardiovascular pregnancy complications, Scimitar Syndrome, cardiovascular syphilis, and cardiovascular tuberculosis.

Arrhythmias include sinus arrhythmia, atrial fibrillation, atrial flutter, bradycardia, extrasystole, Adams-Stokes Syndrome, bundle-branch block, sinoatrial block, long QT syndrome, parasystole, Lown-Ganong-Levine Syndrome, Mahaim-type pre-excitation 30 syndrome, Wolff-Parkinson-White syndrome, sick sinus syndrome, tachycardias, and ventricular fibrillation. Tachycardias include paroxysmal tachycardia, supraventricular tachycardia, accelerated idioventricular rhythm, atrioventricular nodal reentry tachycardia,

ectopic atrial tachycardia, ectopic junctional tachycardia, sinoatrial nodal reentry tachycardia, sinus tachycardia, Torsades de Pointes, and ventricular tachycardia.

Heart valve disease include aortic valve insufficiency, aortic valve stenosis, heart murmurs, aortic valve prolapse, mitral valve prolapse, tricuspid valve prolapse, mitral valve insufficiency, mitral valve stenosis, pulmonary atresia, pulmonary valve insufficiency, pulmonary valve stenosis, tricuspid atresia, tricuspid valve insufficiency, and tricuspid valve stenosis.

Myocardial diseases include alcoholic cardiomyopathy, congestive cardiomyopathy, hypertrophic cardiomyopathy, aortic subvalvular stenosis, pulmonary subvalvular stenosis, restrictive cardiomyopathy, Chagas cardiomyopathy, endocardial fibroelastosis, endomyocardial fibrosis, Kearns Syndrome, myocardial reperfusion injury, and myocarditis.

Myocardial ischemias include coronary disease, such as angina pectoris, coronary aneurysm, coronary arteriosclerosis, coronary thrombosis, coronary vasospasm, myocardial infarction and myocardial stunning.

Cardiovascular diseases also include vascular diseases such as aneurysms, angiodyplasia, angiomatosis, bacillary angiomatosis, Hippel-Lindau Disease, Klippel-Trenaunay-Weber Syndrome, Sturge-Weber Syndrome, angioneurotic edema, aortic diseases, Takayasu's Arteritis, aortitis, Leriche's Syndrome, arterial occlusive diseases, arteritis, enarteritis, polyarteritis nodosa, cerebrovascular disorders, diabetic angiopathies, diabetic retinopathy, embolisms, thrombosis, erythromelalgia, hemorrhoids, hepatic veno-occlusive disease, hypertension, hypotension, ischemia, peripheral vascular diseases, phlebitis, pulmonary veno-occlusive disease, Raynaud's disease, CREST syndrome, retinal vein occlusion, Scimitar syndrome, superior vena cava syndrome, telangiectasia, atacia telangiectasia, hereditary hemorrhagic telangiectasia, varicocele, varicose veins, varicose ulcer, vasculitis, and venous insufficiency.

Aneurysms include dissecting aneurysms, false aneurysms, infected aneurysms, ruptured aneurysms, aortic aneurysms, cerebral aneurysms, coronary aneurysms, heart aneurysms, and iliac aneurysms.

Arterial occlusive diseases include arteriosclerosis, intermittent claudication, carotid stenosis, fibromuscular dysplasias, mesenteric vascular occlusion, Moyamoya disease, renal artery obstruction, retinal artery occlusion, and thromboangiitis obliterans.

Cerebrovascular disorders include carotid artery diseases, cerebral amyloid angiopathy, cerebral aneurysm, cerebral anoxia, cerebral arteriosclerosis, cerebral arteriovenous malformation, cerebral artery diseases, cerebral embolism and thrombosis, carotid artery thrombosis, sinus thrombosis, Wallenberg's syndrome, cerebral hemorrhage, epidural hematoma, subdural hematoma, subarachnoid hemorrhage, cerebral infarction, cerebral ischemia (including transient), subclavian steal syndrome, periventricular leukomalacia, vascular headache, cluster headache, migraine, and vertebrobasilar insufficiency.

Embolisms include air embolisms, amniotic fluid embolisms, cholesterol embolisms, blue toe syndrome, fat embolisms, pulmonary embolisms, and thromboembolisms. Thrombosis include coronary thrombosis, hepatic vein thrombosis, retinal vein occlusion, carotid artery thrombosis, sinus thrombosis, Wallenberg's syndrome, and thrombophlebitis.

Ischemia includes cerebral ischemia, ischemic colitis, compartment syndromes, anterior compartment syndrome, myocardial ischemia, reperfusion injuries, and peripheral limb ischemia. Vasculitis includes aortitis, arteritis, Behcet's Syndrome, Churg-Strauss Syndrome, mucocutaneous lymph node syndrome, thromboangiitis obliterans, hypersensitivity vasculitis, Schoenlein-Henoch purpura, allergic cutaneous vasculitis, and Wegener's granulomatosis.

Polynucleotides or polypeptides, or agonists or antagonists of the present invention, are especially effective for the treatment of critical limb ischemia and coronary disease.

Polypeptides may be administered using any method known in the art, including, but not limited to, direct needle injection at the delivery site, intravenous injection, topical administration, catheter infusion, biolistic injectors, particle accelerators, gelfoam sponge depots, other commercially available depot materials, osmotic pumps, oral or suppository solid pharmaceutical formulations, decanting or topical applications during surgery, aerosol delivery. Such methods are known in the art. Polypeptides may be administered as part of a Therapeutic, described in more detail below. Methods of delivering polynucleotides are described in more detail herein.

Anti-Angiogenesis Activity

The naturally occurring balance between endogenous stimulators and inhibitors of angiogenesis is one in which inhibitory influences predominate. Rastinejad *et al.*, *Cell*

56:345-355 (1989). In those rare instances in which neovascularization occurs under normal physiological conditions, such as wound healing, organ regeneration, embryonic development, and female reproductive processes, angiogenesis is stringently regulated and spatially and temporally delimited. Under conditions of pathological angiogenesis such as that characterizing solid tumor growth, these regulatory controls fail. Unregulated angiogenesis becomes pathologic and sustains progression of many neoplastic and non-neoplastic diseases. A number of serious diseases are dominated by abnormal neovascularization including solid tumor growth and metastases, arthritis, some types of eye disorders, and psoriasis. See, e.g., reviews by Moses *et al.*, *Biotech.* 9:630-634 (1991); Folkman *et al.*, *N. Engl. J. Med.*, 333:1757-1763 (1995); Auerbach *et al.*, *J. Microvasc. Res.* 29:401-411 (1985); Folkman, *Advances in Cancer Research*, eds. Klein and Weinhouse, Academic Press, New York, pp. 175-203 (1985); Patz, *Am. J. Ophthalmol.* 94:715-743 (1982); and Folkman *et al.*, *Science* 221:719-725 (1983). In a number of pathological conditions, the process of angiogenesis contributes to the disease state. For example, significant data have accumulated which suggest that the growth of solid tumors is dependent on angiogenesis. Folkman and Klagsbrun, *Science* 235:442-447 (1987).

The present invention provides for treatment of diseases or disorders associated with neovascularization by administration of the polynucleotides and/or polypeptides of the invention, as well as agonists or antagonists of the present invention. Malignant and metastatic conditions which can be treated with the polynucleotides and polypeptides, or agonists or antagonists of the invention include, but are not limited to, malignancies, solid tumors, and cancers described herein and otherwise known in the art (for a review of such disorders, see Fishman *et al.*, *Medicine*, 2d Ed., J. B. Lippincott Co., Philadelphia (1985)). Thus, the present invention provides a method of treating an angiogenesis-related disease and/or disorder, comprising administering to an individual in need thereof a therapeutically effective amount of a polynucleotide, polypeptide, antagonist and/or agonist of the invention. For example, polynucleotides, polypeptides, antagonists and/or agonists may be utilized in a variety of additional methods in order to therapeutically treat a cancer or tumor. Cancers which may be treated with polynucleotides, polypeptides, antagonists and/or agonists include, but are not limited to solid tumors, including prostate, lung, breast, ovarian, stomach, pancreas, larynx, esophagus, testes, liver, parotid, biliary tract, colon, rectum, cervix, uterus, endometrium, kidney, bladder, thyroid cancer; primary tumors and metastases;

melanomas; glioblastoma; Kaposi's sarcoma; leiomyosarcoma; non- small cell lung cancer; colorectal cancer; advanced malignancies; and blood born tumors such as leukemias. For example, polynucleotides, polypeptides, antagonists and/or agonists may be delivered topically, in order to treat cancers such as skin cancer, head and neck tumors, breast tumors,
5 and Kaposi's sarcoma.

Within yet other aspects, polynucleotides, polypeptides, antagonists and/or agonists may be utilized to treat superficial forms of bladder cancer by, for example, intravesical administration. Polynucleotides, polypeptides, antagonists and/or agonists may be delivered directly into the tumor, or near the tumor site, via injection or a catheter. Of course, as the
10 artisan of ordinary skill will appreciate, the appropriate mode of administration will vary according to the cancer to be treated. Other modes of delivery are discussed herein.

Polynucleotides, polypeptides, antagonists and/or agonists may be useful in treating other disorders, besides cancers, which involve angiogenesis. These disorders include, but are not limited to: benign tumors, for example hemangiomas, acoustic neuromas,
15 neurofibromas, trachomas, and pyogenic granulomas; arteriosclerotic plaques; ocular angiogenic diseases, for example, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, retinoblastoma, uveitis and Pterygia (abnormal blood vessel growth) of the eye; rheumatoid arthritis; psoriasis; delayed wound healing; endometriosis; vasculogenesis; granulations;
20 hypertrophic scars (keloids); nonunion fractures; scleroderma; trachoma; vascular adhesions; myocardial angiogenesis; coronary collaterals; cerebral collaterals; arteriovenous malformations; ischemic limb angiogenesis; Osler-Webber Syndrome; plaque neovascularization; telangiectasia; hemophilic joints; angiofibroma; fibromuscular dysplasia; wound granulation; Crohn's disease; and atherosclerosis.

For example, within one aspect of the present invention methods are provided for treating hypertrophic scars and keloids, comprising the step of administering a polynucleotide, polypeptide, antagonist and/or agonist of the invention to a hypertrophic scar or keloid.
25

Within one embodiment of the present invention polynucleotides, polypeptides, antagonists and/or agonists are directly injected into a hypertrophic scar or keloid, in order to prevent the progression of these lesions. This therapy is of particular value in the prophylactic treatment of conditions which are known to result in the development of
30

hypertrophic scars and keloids (e.g., burns), and is preferably initiated after the proliferative phase has had time to progress (approximately 14 days after the initial injury), but before hypertrophic scar or keloid development. As noted above, the present invention also provides methods for treating neovascular diseases of the eye, including for example, corneal
5 neovascularization, neovascular glaucoma, proliferative diabetic retinopathy, retrolental fibroplasia and macular degeneration.

Moreover, Ocular disorders associated with neovascularization which can be treated with the polynucleotides and polypeptides of the present invention (including agonists and/or antagonists) include, but are not limited to: neovascular glaucoma, diabetic retinopathy,
10 retinoblastoma, retrolental fibroplasia, uveitis, retinopathy of prematurity macular degeneration, corneal graft neovascularization, as well as other eye inflammatory diseases, ocular tumors and diseases associated with choroidal or iris neovascularization. See, e.g., reviews by Waltman *et al.*, *Am. J. Ophthalm.* 85:704-710 (1978) and Gartner *et al.*, *Surv. Ophthalm.* 22:291-312 (1978).

15 Thus, within one aspect of the present invention methods are provided for treating neovascular diseases of the eye such as corneal neovascularization (including corneal graft neovascularization), comprising the step of administering to a patient a therapeutically effective amount of a compound (as described above) to the cornea, such that the formation of blood vessels is inhibited. Briefly, the cornea is a tissue which normally lacks blood
20 vessels. In certain pathological conditions however, capillaries may extend into the cornea from the pericorneal vascular plexus of the limbus. When the cornea becomes vascularized, it also becomes clouded, resulting in a decline in the patient's visual acuity. Visual loss may become complete if the cornea completely opacitates. A wide variety of disorders can result in corneal neovascularization, including for example, corneal infections (e.g., trachoma,
25 herpes simplex keratitis, leishmaniasis and onchocerciasis), immunological processes (e.g., graft rejection and Stevens-Johnson's syndrome), alkali burns, trauma, inflammation (of any cause), toxic and nutritional deficiency states, and as a complication of wearing contact lenses.

Within particularly preferred embodiments of the invention, may be prepared for
30 topical administration in saline (combined with any of the preservatives and antimicrobial agents commonly used in ocular preparations), and administered in eyedrop form. The solution or suspension may be prepared in its pure form and administered several times daily.

Alternatively, anti-angiogenic compositions, prepared as described above, may also be administered directly to the cornea. Within preferred embodiments, the anti-angiogenic composition is prepared with a muco-adhesive polymer which binds to cornea. Within further embodiments, the anti-angiogenic factors or anti-angiogenic compositions may be
5 utilized as an adjunct to conventional steroid therapy. Topical therapy may also be useful prophylactically in corneal lesions which are known to have a high probability of inducing an angiogenic response (such as chemical burns). In these instances the treatment, likely in combination with steroids, may be instituted immediately to help prevent subsequent complications.

10 Within other embodiments, the compounds described above may be injected directly into the corneal stroma by an ophthalmologist under microscopic guidance. The preferred site of injection may vary with the morphology of the individual lesion, but the goal of the administration would be to place the composition at the advancing front of the vasculature (i.e., interspersed between the blood vessels and the normal cornea). In most cases this
15 would involve perilimbic corneal injection to "protect" the cornea from the advancing blood vessels. This method may also be utilized shortly after a corneal insult in order to prophylactically prevent corneal neovascularization. In this situation the material could be injected in the perilimbic cornea interspersed between the corneal lesion and its undesired potential limbic blood supply. Such methods may also be utilized in a similar fashion to
20 prevent capillary invasion of transplanted corneas. In a sustained-release form injections might only be required 2-3 times per year. A steroid could also be added to the injection solution to reduce inflammation resulting from the injection itself.

Within another aspect of the present invention, methods are provided for treating neovascular glaucoma, comprising the step of administering to a patient a therapeutically
25 effective amount of a polynucleotide, polypeptide, antagonist and/or agonist to the eye, such that the formation of blood vessels is inhibited. In one embodiment, the compound may be administered topically to the eye in order to treat early forms of neovascular glaucoma. Within other embodiments, the compound may be implanted by injection into the region of the anterior chamber angle. Within other embodiments, the compound may also be placed in
30 any location such that the compound is continuously released into the aqueous humor. Within another aspect of the present invention, methods are provided for treating proliferative diabetic retinopathy, comprising the step of administering to a patient a

therapeutically effective amount of a polynucleotide, polypeptide, antagonist and/or agonist to the eyes, such that the formation of blood vessels is inhibited.

Within particularly preferred embodiments of the invention, proliferative diabetic retinopathy may be treated by injection into the aqueous humor or the vitreous, in order to
5 increase the local concentration of the polynucleotide, polypeptide, antagonist and/or agonist in the retina. Preferably, this treatment should be initiated prior to the acquisition of severe disease requiring photocoagulation.

Within another aspect of the present invention, methods are provided for treating retrolental fibroplasia, comprising the step of administering to a patient a therapeutically
10 effective amount of a polynucleotide, polypeptide, antagonist and/or agonist to the eye, such that the formation of blood vessels is inhibited. The compound may be administered topically, via intravitreal injection and/or via intraocular implants.

Additionally, disorders which can be treated with the polynucleotides, polypeptides, agonists and/or antagonists include, but are not limited to, hemangioma, arthritis, psoriasis, angiofibroma, atherosclerotic plaques, delayed wound healing, granulations, hemophilic
15 joints, hypertrophic scars, nonunion fractures, Osler-Weber syndrome, pyogenic granuloma, scleroderma, trachoma, and vascular adhesions.

Moreover, disorders and/or states, which can be treated with the polynucleotides, polypeptides, agonists and/or antagonists include, but are not limited to, solid
20 tumors, blood born tumors such as leukemias, tumor metastasis, Kaposi's sarcoma, benign tumors, for example hemangiomas, acoustic neuromas, neurofibromas, trachomas, and pyogenic granulomas, rheumatoid arthritis, psoriasis, ocular angiogenic diseases, for example, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, retinoblastoma, and
25 uveitis, delayed wound healing, endometriosis, vasculogenesis, granulations, hypertrophic scars (keloids), nonunion fractures, scleroderma, trachoma, vascular adhesions, myocardial angiogenesis, coronary collaterals, cerebral collaterals, arteriovenous malformations, ischemic limb angiogenesis, Osler-Webber Syndrome, plaque neovascularization, telangiectasia, hemophilic joints, angiofibroma fibromuscular dysplasia, wound granulation,
30 Crohn's disease, atherosclerosis, birth control agent by preventing vascularization required for embryo implantation controlling menstruation, diseases that have angiogenesis as a

pathologic consequence such as cat scratch disease (*Rochela minalia quintosa*), ulcers (*Helicobacter pylori*), Bartonellosis and bacillary angiomatosis.

In one aspect of the birth control method, an amount of the compound sufficient to block embryo implantation is administered before or after intercourse and fertilization have occurred, thus providing an effective method of birth control, possibly a "morning after" method. Polynucleotides, polypeptides, agonists and/or agonists may also be used in controlling menstruation or administered as either a peritoneal lavage fluid or for peritoneal implantation in the treatment of endometriosis.

Polynucleotides, polypeptides, agonists and/or agonists of the present invention may be incorporated into surgical sutures in order to prevent stitch granulomas.

Polynucleotides, polypeptides, agonists and/or agonists may be utilized in a wide variety of surgical procedures. For example, within one aspect of the present invention a compositions (in the form of, for example, a spray or film) may be utilized to coat or spray an area prior to removal of a tumor, in order to isolate normal surrounding tissues from malignant tissue, and/or to prevent the spread of disease to surrounding tissues. Within other aspects of the present invention, compositions (e.g., in the form of a spray) may be delivered via endoscopic procedures in order to coat tumors, or inhibit angiogenesis in a desired locale. Within yet other aspects of the present invention, surgical meshes which have been coated with anti-angiogenic compositions of the present invention may be utilized in any procedure wherein a surgical mesh might be utilized. For example, within one embodiment of the invention a surgical mesh laden with an anti-angiogenic composition may be utilized during abdominal cancer resection surgery (e.g., subsequent to colon resection) in order to provide support to the structure, and to release an amount of the anti-angiogenic factor.

Within further aspects of the present invention, methods are provided for treating tumor excision sites, comprising administering a polynucleotide, polypeptide, agonist and/or agonist to the resection margins of a tumor subsequent to excision, such that the local recurrence of cancer and the formation of new blood vessels at the site is inhibited. Within one embodiment of the invention, the anti-angiogenic compound is administered directly to the tumor excision site (e.g., applied by swabbing, brushing or otherwise coating the resection margins of the tumor with the anti-angiogenic compound). Alternatively, the anti-angiogenic compounds may be incorporated into known surgical pastes prior to administration. Within particularly preferred embodiments of the invention, the anti-

angiogenic compounds are applied after hepatic resections for malignancy, and after neurosurgical operations.

Within one aspect of the present invention, polynucleotides, polypeptides, agonists and/or agonists may be administered to the resection margin of a wide variety of tumors, including for example, breast, colon, brain and hepatic tumors. For example, within one embodiment of the invention, anti-angiogenic compounds may be administered to the site of a neurological tumor subsequent to excision, such that the formation of new blood vessels at the site are inhibited.

The polynucleotides, polypeptides, agonists and/or agonists of the present invention may also be administered along with other anti-angiogenic factors. Representative examples of other anti-angiogenic factors include: Anti-Invasive Factor, retinoic acid and derivatives thereof, paclitaxel, Suramin, Tissue Inhibitor of Metalloproteinase-1, Tissue Inhibitor of Metalloproteinase-2, Plasminogen Activator Inhibitor-1, Plasminogen Activator Inhibitor-2, and various forms of the lighter "d group" transition metals.

Lighter "d group" transition metals include, for example, vanadium, molybdenum, tungsten, titanium, niobium, and tantalum species. Such transition metal species may form transition metal complexes. Suitable complexes of the above-mentioned transition metal species include oxo transition metal complexes.

Representative examples of vanadium complexes include oxo vanadium complexes such as vanadate and vanadyl complexes. Suitable vanadate complexes include metavanadate and orthovanadate complexes such as, for example, ammonium metavanadate, sodium metavanadate, and sodium orthovanadate. Suitable vanadyl complexes include, for example, vanadyl acetylacetonate and vanadyl sulfate including vanadyl sulfate hydrates such as vanadyl sulfate mono- and trihydrates.

Representative examples of tungsten and molybdenum complexes also include oxo complexes. Suitable oxo tungsten complexes include tungstate and tungsten oxide complexes. Suitable tungstate complexes include ammonium tungstate, calcium tungstate, sodium tungstate dihydrate, and tungstic acid. Suitable tungsten oxides include tungsten (IV) oxide and tungsten (VI) oxide. Suitable oxo molybdenum complexes include molybdate, molybdenum oxide, and molybdenyl complexes. Suitable molybdate complexes include ammonium molybdate and its hydrates, sodium molybdate and its hydrates, and potassium molybdate and its hydrates. Suitable molybdenum oxides include molybdenum (VI) oxide,

molybdenum (VI) oxide, and molybdic acid. Suitable molybdenyl complexes include, for example, molybdenyl acetylacetonate. Other suitable tungsten and molybdenum complexes include hydroxo derivatives derived from, for example, glycerol, tartaric acid, and sugars.

A wide variety of other anti-angiogenic factors may also be utilized within the context
5 of the present invention. Representative examples include platelet factor 4; protamine sulphate; sulphated chitin derivatives (prepared from queen crab shells), (Murata et al., Cancer Res. 51:22-26, 1991); Sulphated Polysaccharide Peptidoglycan Complex (SP- PG) (the function of this compound may be enhanced by the presence of steroids such as estrogen, and tamoxifen citrate); Staurosporine; modulators of matrix metabolism, including for
10 example, proline analogs, cishydroxyproline, d,L-3,4-dehydroproline, Thiaproline, alpha,alpha-dipyridyl, aminopropionitrile fumarate; 4-propyl-5-(4-pyridinyl)-2(3H)-oxazolone; Methotrexate; Mitoxantrone; Heparin; Interferons; 2 Macroglobulin-serum; ChIMP-3 (Pavloff et al., J. Bio. Chem. 267:17321-17326, 1992); Chymostatin (Tomkinson et al., Biochem J. 286:475-480, 1992); Cyclodextrin Tetradecasulfate; Eponemycin;
15 Camptothecin; Fumagillin (Ingber et al., Nature 348:555-557, 1990); Gold Sodium Thiomalate ("GST"; Matsubara and Ziff, J. Clin. Invest. 79:1440-1446, 1987); anticollagenase-serum; alpha2-antiplasmin (Holmes et al., J. Biol. Chem. 262(4):1659-1664, 1987); Bisantrene (National Cancer Institute); Lobenzarit disodium (N-(2)-carboxyphenyl-4-chloroanthronilic acid disodium or "CCA"; Takeuchi et al., Agents Actions 36:312-316,
20 1992); Thalidomide; Angostatic steroid; AGM-1470; carboxynaminolmidazole; and metalloproteinase inhibitors such as BB94.

Diseases at the Cellular Level

Diseases associated with increased cell survival or the inhibition of apoptosis that
25 could be treated or detected by polynucleotides or polypeptides, as well as antagonists or agonists of the present invention, include cancers (such as follicular lymphomas, carcinomas with p53 mutations, and hormone-dependent tumors, including, but not limited to colon cancer, cardiac tumors, pancreatic cancer, melanoma, retinoblastoma, glioblastoma, lung cancer, intestinal cancer, testicular cancer, stomach cancer, neuroblastoma, myxoma, myoma,
30 lymphoma, endothelioma, osteoblastoma, osteoclastoma, osteosarcoma, chondrosarcoma, adenoma, breast cancer, prostate cancer, Kaposi's sarcoma and ovarian cancer); autoimmune disorders (such as, multiple sclerosis, Sjogren's syndrome, Hashimoto's thyroiditis, biliary

cirrhosis, Behcet's disease, Crohn's disease, polymyositis, systemic lupus erythematosus and immune-related glomerulonephritis and rheumatoid arthritis) and viral infections (such as herpes viruses, pox viruses and adenoviruses), inflammation, graft v. host disease, acute graft rejection, and chronic graft rejection. In preferred embodiments, polynucleotides, polypeptides, and/or antagonists of the invention are used to inhibit growth, progression, and/or metasis of cancers, in particular those listed above.

Additional diseases or conditions associated with increased cell survival that could be treated or detected by polynucleotides or polypeptides, or agonists or antagonists of the present invention include, but are not limited to, progression, and/or metastases of malignancies and related disorders such as leukemia (including acute leukemias (e.g., acute lymphocytic leukemia, acute myelocytic leukemia (including myeloblastic, promyelocytic, myelomonocytic, monocytic, and erythroleukemia)) and chronic leukemias (e.g., chronic myelocytic (granulocytic) leukemia and chronic lymphocytic leukemia)), polycythemia vera, lymphomas (e.g., Hodgkin's disease and non-Hodgkin's disease), multiple myeloma, Waldenstrom's macroglobulinemia, heavy chain disease, and solid tumors including, but not limited to, sarcomas and carcinomas such as fibrosarcoma, myxosarcoma, liposarcoma, chondrosarcoma, osteogenic sarcoma, chordoma, angiosarcoma, endotheliosarcoma, lymphangiosarcoma, lymphangioendotheliosarcoma, synovioma, mesothelioma, Ewing's tumor, leiomyosarcoma, rhabdomyosarcoma, colon carcinoma, pancreatic cancer, breast cancer, ovarian cancer, prostate cancer, squamous cell carcinoma, basal cell carcinoma, adenocarcinoma, sweat gland carcinoma, sebaceous gland carcinoma, papillary carcinoma, papillary adenocarcinomas, cystadenocarcinoma, medullary carcinoma, bronchogenic carcinoma, renal cell carcinoma, hepatoma, bile duct carcinoma, choriocarcinoma, seminoma, embryonal carcinoma, Wilm's tumor, cervical cancer, testicular tumor, lung carcinoma, small cell lung carcinoma, bladder carcinoma, epithelial carcinoma, glioma, astrocytoma, medulloblastoma, craniopharyngioma, ependymoma, pinealoma, hemangioblastoma, acoustic neuroma, oligodendroglioma, menangioma, melanoma, neuroblastoma, and retinoblastoma.

Diseases associated with increased apoptosis that could be treated or detected by polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, include AIDS; neurodegenerative disorders (such as Alzheimer's disease, Parkinson's disease, Amyotrophic lateral sclerosis, Retinitis pigmentosa, Cerebellar degeneration and brain tumor

or prior associated disease); autoimmune disorders (such as, multiple sclerosis, Sjogren's syndrome, Hashimoto's thyroiditis, biliary cirrhosis, Behcet's disease, Crohn's disease, polymyositis, systemic lupus erythematosus and immune-related glomerulonephritis and rheumatoid arthritis) myelodysplastic syndromes (such as aplastic anemia), graft v. host disease, ischemic injury (such as that caused by myocardial infarction, stroke and reperfusion injury), liver injury (e.g., hepatitis related liver injury, ischemia/reperfusion injury, cholestasis (bile duct injury) and liver cancer); toxin-induced liver disease (such as that caused by alcohol), septic shock, cachexia and anorexia.

10 **Wound Healing and Epithelial Cell Proliferation**

In accordance with yet a further aspect of the present invention, there is provided a process for utilizing polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, for therapeutic purposes, for example, to stimulate epithelial cell proliferation and basal keratinocytes for the purpose of wound healing, and to stimulate hair follicle production and healing of dermal wounds. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, may be clinically useful in stimulating wound healing including surgical wounds, excisional wounds, deep wounds involving damage of the dermis and epidermis, eye tissue wounds, dental tissue wounds, oral cavity wounds, diabetic ulcers, dermal ulcers, cubitus ulcers, arterial ulcers, venous stasis ulcers, burns resulting from heat exposure or chemicals, and other abnormal wound healing conditions such as uremia, malnutrition, vitamin deficiencies and complications associated with systemic treatment with steroids, radiation therapy and antineoplastic drugs and antimetabolites. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to promote dermal reestablishment subsequent to dermal loss

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to increase the adherence of skin grafts to a wound bed and to stimulate re-epithelialization from the wound bed. The following are types of grafts that polynucleotides or polypeptides, agonists or antagonists of the present invention, could be used to increase adherence to a wound bed: autografts, artificial skin, allografts, autodermic graft, autoepidermic grafts, avacular grafts, Blair-Brown grafts, bone graft, brephoplastic grafts, cutis graft, delayed graft, dermic graft, epidermic graft, fascia graft, full thickness

graft, heterologous graft, xenograft, homologous graft, hyperplastic graft, lamellar graft, mesh graft, mucosal graft, Ollier-Thiersch graft, omentoplastic graft, patch graft, pedicle graft, penetrating graft, split skin graft, thick split graft. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, can be used to promote skin strength and to
5 improve the appearance of aged skin.

It is believed that polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, will also produce changes in hepatocyte proliferation, and epithelial cell proliferation in the lung, breast, pancreas, stomach, small intestine, and large intestine. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention,
10 could promote proliferation of epithelial cells such as sebocytes, hair follicles, hepatocytes, type II pneumocytes, mucin-producing goblet cells, and other epithelial cells and their progenitors contained within the skin, lung, liver, and gastrointestinal tract. Polynucleotides or polypeptides, agonists or antagonists of the present invention, may promote proliferation of endothelial cells, keratinocytes, and basal keratinocytes.

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could also be used to reduce the side effects of gut toxicity that result from radiation, chemotherapy treatments or viral infections. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, may have a cytoprotective effect on the small intestine mucosa. Polynucleotides or polypeptides, as well as agonists or
20 antagonists of the present invention, may also stimulate healing of mucositis (mouth ulcers) that result from chemotherapy and viral infections.

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could further be used in full regeneration of skin in full and partial thickness skin defects, including burns, (i.e., repopulation of hair follicles, sweat glands, and sebaceous
25 glands), treatment of other skin defects such as psoriasis. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to treat epidermolysis bullosa, a defect in adherence of the epidermis to the underlying dermis which results in frequent, open and painful blisters by accelerating reepithelialization of these lesions. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention,
30 could also be used to treat gastric and duodenal ulcers and help heal by scar formation of the mucosal lining and regeneration of glandular mucosa and duodenal mucosal lining more rapidly. Inflammatory bowel diseases, such as Crohn's disease and ulcerative colitis, are

diseases which result in destruction of the mucosal surface of the small or large intestine, respectively. Thus, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to promote the resurfacing of the mucosal surface to aid more rapid healing and to prevent progression of inflammatory bowel disease. Treatment with polynucleotides or polypeptides, agonists or antagonists of the present invention, is expected to have a significant effect on the production of mucus throughout the gastrointestinal tract and could be used to protect the intestinal mucosa from injurious substances that are ingested or following surgery. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to treat diseases associated with the under expression.

Moreover, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to prevent and heal damage to the lungs due to various pathological states. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, which could stimulate proliferation and differentiation and promote the repair of alveoli and bronchiolar epithelium to prevent or treat acute or chronic lung damage. For example, emphysema, which results in the progressive loss of alveoli, and inhalation injuries, i.e., resulting from smoke inhalation and burns, that cause necrosis of the bronchiolar epithelium and alveoli could be effectively treated using polynucleotides or polypeptides, agonists or antagonists of the present invention. Also, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to stimulate the proliferation of and differentiation of type II pneumocytes, which may help treat or prevent disease such as hyaline membrane diseases, such as infant respiratory distress syndrome and bronchopulmonary dysplasia, in premature infants.

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could stimulate the proliferation and differentiation of hepatocytes and, thus, could be used to alleviate or treat liver diseases and pathologies such as fulminant liver failure caused by cirrhosis, liver damage caused by viral hepatitis and toxic substances (i.e., acetaminophen, carbon tetrachloride and other hepatotoxins known in the art).

In addition, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used to treat or prevent the onset of diabetes mellitus. In patients with newly diagnosed Types I and II diabetes, where some islet cell function remains, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention,

could be used to maintain the islet function so as to alleviate, delay or prevent permanent manifestation of the disease. Also, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, could be used as an auxiliary in islet cell transplantation to improve or promote islet cell function.

5

Neurological Diseases

In accordance with yet a further aspect of the present invention, there is provided a process for utilizing polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, for therapeutic purposes, for example, to stimulate neurological cell proliferation and/or differentiation. Therefore, polynucleotides, polypeptides, agonists and/or antagonists of the invention may be used to treat and/or detect neurologic diseases. Moreover, polynucleotides or polypeptides, or agonists or antagonists of the invention, can be used as a marker or detector of a particular nervous system disease or disorder.

Examples of neurologic diseases which can be treated or detected with polynucleotides, polypeptides, agonists, and/or antagonists of the present invention include brain diseases, such as metabolic brain diseases which includes phenylketonuria such as maternal phenylketonuria, pyruvate carboxylase deficiency, pyruvate dehydrogenase complex deficiency, Wernicke's Encephalopathy, brain edema, brain neoplasms such as cerebellar neoplasms which include infratentorial neoplasms, cerebral ventricle neoplasms such as choroid plexus neoplasms, hypothalamic neoplasms, supratentorial neoplasms, canavan disease, cerebellar diseases such as cerebellar ataxia which include spinocerebellar degeneration such as ataxia telangiectasia, cerebellar dyssynergia, Friederich's Ataxia, Machado-Joseph Disease, olivopontocerebellar atrophy, cerebellar neoplasms such as infratentorial neoplasms, diffuse cerebral sclerosis such as encephalitis periaxialis, globoid cell leukodystrophy, metachromatic leukodystrophy and subacute sclerosing panencephalitis, cerebrovascular disorders (such as carotid artery diseases which include carotid artery thrombosis, carotid stenosis and Moyamoya Disease, cerebral amyloid angiopathy, cerebral aneurysm, cerebral anoxia, cerebral arteriosclerosis, cerebral arteriovenous malformations, cerebral artery diseases, cerebral embolism and thrombosis such as carotid artery thrombosis, sinus thrombosis and Wallenberg's Syndrome, cerebral hemorrhage such as epidural hematoma, subdural hematoma and subarachnoid hemorrhage, cerebral infarction, cerebral ischemia such as transient cerebral ischemia, Subclavian Steal Syndrome and vertebrobasilar

insufficiency, vascular dementia such as multi-infarct dementia, periventricular leukomalacia, vascular headache such as cluster headache, migraine, dementia such as AIDS Dementia Complex, presenile dementia such as Alzheimer's Disease and Creutzfeldt-Jakob Syndrome, senile dementia such as Alzheimer's Disease and progressive supranuclear palsy, vascular

5 dementia such as multi-infarct dementia, encephalitis which include encephalitis periaxialis, viral encephalitis such as epidemic encephalitis, Japanese Encephalitis, St. Louis Encephalitis, tick-borne encephalitis and West Nile Fever, acute disseminated encephalomyelitis, meningoencephalitis such as uveomeningoencephalitic syndrome, Postencephalitic Parkinson Disease and subacute sclerosing panencephalitis,

10 encephalomalacia such as periventricular leukomalacia, epilepsy such as generalized epilepsy which includes infantile spasms, absence epilepsy, myoclonic epilepsy which includes MERRF Syndrome, tonic-clonic epilepsy, partial epilepsy such as complex partial epilepsy, frontal lobe epilepsy and temporal lobe epilepsy, post-traumatic epilepsy, status epilepticus such as Epilepsia Partialis Continua, Hallervorden-Spatz Syndrome, hydrocephalus such as

15 Dandy-Walker Syndrome and normal pressure hydrocephalus, hypothalamic diseases such as hypothalamic neoplasms, cerebral malaria, narcolepsy which includes cataplexy, bulbar poliomyelitis, cerebri pseudotumor, Rett Syndrome, Reye's Syndrome, thalamic diseases, cerebral toxoplasmosis, intracranial tuberculoma and Zellweger Syndrome, central nervous system infections such as AIDS Dementia Complex, Brain Abscess, subdural empyema,

20 encephalomyelitis such as Equine Encephalomyelitis, Venezuelan Equine Encephalomyelitis, Necrotizing Hemorrhagic Encephalomyelitis, Visna, cerebral malaria, meningitis such as arachnoiditis, aseptic meningitis such as viral meningitis which includes lymphocytic choriomeningitis. Bacterial meningitis which includes Haemophilus Meningitis, Listeria Meningitis, Meningococcal Meningitis such as Waterhouse-Friderichsen Syndrome,

25 Pneumococcal Meningitis and meningeal tuberculosis, fungal meningitis such as Cryptococcal Meningitis, subdural effusion, meningoencephalitis such as uveomeningoencephalitic syndrome, myelitis such as transverse myelitis, neurosyphilis such as tabes dorsalis, poliomyelitis which includes bulbar poliomyelitis and postpoliomyelitis syndrome, prion diseases (such as Creutzfeldt-Jakob Syndrome, Bovine Spongiform

30 Encephalopathy, Gerstmann-Straussler Syndrome, Kuru, Scrapie) cerebral toxoplasmosis, central nervous system neoplasms such as brain neoplasms that include cerebellar neoplasms such as infratentorial neoplasms, cerebral ventricle neoplasms such as choroid

plexus neoplasms, hypothalamic neoplasms and supratentorial neoplasms, meningeal neoplasms, spinal cord neoplasms which include epidural neoplasms, demyelinating diseases such as Canavan Diseases, diffuse cerebral sclerolosis which includes adrenoleukodystrophy, encephalitis periaxialis, globoid cell leukodystrophy, diffuse cerebral sclerosis such as metachromatic leukodystrophy, allergic encephalomyelitis, necrotizing hemorrhagic encephalomyelitis, progressive multifocal leukoencephalopathy, multiple sclerosis, central pontine myelinolysis, transverse myelitis, neuromyelitis optica, Scrapie, Swayback, Chronic Fatigue Syndrome, Visna, High Pressure Nervous Syndrome, Meningism, spinal cord diseases such as amyotonia congenita, amyotrophic lateral sclerosis, spinal muscular atrophy such as Werdnig-Hoffmann Disease, spinal cord compression, spinal cord neoplasms such as epidural neoplasms, syringomyelia, Tabes Dorsalis, Stiff-Man Syndrome, mental retardation such as Angelman Syndrome, Cri-du-Chat Syndrome, De Lange's Syndrome, Down Syndrome, Gangliosidoses such as gangliosidoses G(M1), Sandhoff Disease, Tay-Sachs Disease, Hartnup Disease, homocystinuria, Laurence-Moon- Biedl Syndrome, Lesch-Nyhan Syndrome, Maple Syrup Urine Disease, mucopolipidosis such as fucosidosis, neuronal ceroid-lipofuscinosis, oculocerebrorenal syndrome, phenylketonuria such as maternal phenylketonuria, Prader-Willi Syndrome, Rett Syndrome, Rubinstein-Taybi Syndrome, Tuberous Sclerosis, WAGR Syndrome, nervous system abnormalities such as holoprosencephaly, neural tube defects such as anencephaly which includes hydrangencephaly, Arnold-Chairi Deformity, encephalocele, meningocele, meningomyelocele, spinal dysraphism such as spina bifida cystica and spina bifida occulta, hereditary motor and sensory neuropathies which include Charcot-Marie Disease, Hereditary optic atrophy, Refsum's Disease, hereditary spastic paraplegia, Werdnig-Hoffmann Disease, Hereditary Sensory and Autonomic Neuropathies such as Congenital Analgesia and Familial Dysautonomia, Neurologic manifestations (such as agnosia that include Gerstmann's Syndrome, Amnesia such as retrograde amnesia, apraxia, neurogenic bladder, cataplexy, communicative disorders such as hearing disorders that includes deafness, partial hearing loss, loudness recruitment and tinnitus, language disorders such as aphasia which include agraphia, anomia, broca aphasia, and Wernicke Aphasia, Dyslexia such as Acquired Dyslexia; language development disorders, speech disorders such as aphasia which includes anomia, broca aphasia and Wernicke Aphasia, articulation disorders, communicative disorders such as speech disorders which include dysarthria, echolalia, mutism and stuttering,

voice disorders such as aphonia and hoarseness, decerebrate state, delirium, fasciculation, hallucinations, meningism, movement disorders such as angelman syndrome, ataxia, athetosis, chorea, dystonia, hypokinesia, muscle hypotonia, myoclonus, tic, torticollis and tremor, muscle hypertonia such as muscle rigidity such as stiff-man syndrome, muscle spasticity, paralysis such as facial paralysis which includes Herpes Zoster Oticus, 5 Gastroparesis, Hemiplegia, ophthalmoplegia such as diplopia, Duane's Syndrome, Horner's Syndrome, Chronic progressive external ophthalmoplegia such as Kearns Syndrome, Bulbar Paralysis, Tropical Spastic Paraparesis, Paraplegia such as Brown-Sequard Syndrome, quadriplegia, respiratory paralysis and vocal cord paralysis, paresis, phantom limb, taste 10 disorders such as ageusia and dysgeusia, vision disorders such as amblyopia, blindness, color vision defects, diplopia, hemianopsia, scotoma and subnormal vision, sleep disorders such as hypersomnia which includes Kleine-Levin Syndrome, insomnia, and somnambulism, spasm such as trismus, unconsciousness such as coma, persistent vegetative state and syncope and vertigo, neuromuscular diseases such as amyotonia congenita, amyotrophic lateral sclerosis, 15 Lambert-Eaton Myasthenic Syndrome, motor neuron disease, muscular atrophy such as spinal muscular atrophy, Charcot-Marie Disease and Werdnig-Hoffmann Disease, Postpoliomyelitis Syndrome, Muscular Dystrophy, Myasthenia Gravis, Myotonia Atrophica, Myotonia Confenita, Nemaline Myopathy, Familial Periodic Paralysis, Multiplex Paramyoclonus, Tropical Spastic Paraparesis and Stiff-Man Syndrome, peripheral nervous 20 system diseases such as acrodynia, amyloid neuropathies, autonomic nervous system diseases such as Adie's Syndrome, Barre-Lieou Syndrome, Familial Dysautonomia, Horner's Syndrome, Reflex Sympathetic Dystrophy and Shy-Drager Syndrome, Cranial Nerve Diseases such as Acoustic Nerve Diseases such as Acoustic Neuroma which includes Neurofibromatosis 2, Facial Nerve Diseases such as Facial Neuralgia, Melkersson-Rosenthal 25 Syndrome, ocular motility disorders which includes amblyopia, nystagmus, oculomotor nerve paralysis, ophthalmoplegia such as Duane's Syndrome, Horner's Syndrome, Chronic Progressive External Ophthalmoplegia which includes Kearns Syndrome, Strabismus such as Esotropia and Exotropia, Oculomotor Nerve Paralysis, Optic Nerve Diseases such as Optic Atrophy which includes Hereditary Optic Atrophy, Optic Disk Drusen, Optic Neuritis such as 30 Neuromyelitis Optica, Papilledema, Trigeminal Neuralgia, Vocal Cord Paralysis, Demyelinating Diseases such as Neuromyelitis Optica and Swayback, Diabetic neuropathies such as diabetic foot, nerve compression syndromes such as carpal tunnel syndrome, tarsal

tunnel syndrome, thoracic outlet syndrome such as cervical rib syndrome, ulnar nerve compression syndrome, neuralgia such as causalgia, cervico-brachial neuralgia, facial neuralgia and trigeminal neuralgia, neuritis such as experimental allergic neuritis, optic neuritis, polyneuritis, polyradiculoneuritis and radiculities such as polyradiculitis, hereditary motor and sensory neuropathies such as Charcot-Marie Disease, Hereditary Optic Atrophy, Refsum's Disease, Hereditary Spastic Paraplegia and Werdnig-Hoffmann Disease, Hereditary Sensory and Autonomic Neuropathies which include Congenital Analgesia and Familial Dysautonomia, POEMS Syndrome, Sciatica, Gustatory Sweating and Tetany).

10 Infectious Disease

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention can be used to treat or detect infectious agents. For example, by increasing the immune response, particularly increasing the proliferation and differentiation of B and/or T cells, infectious diseases may be treated. The immune response may be increased by either enhancing an existing immune response, or by initiating a new immune response. Alternatively, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention may also directly inhibit the infectious agent, without necessarily eliciting an immune response.

Viruses are one example of an infectious agent that can cause disease or symptoms that can be treated or detected by a polynucleotide or polypeptide and/or agonist or antagonist of the present invention. Examples of viruses, include, but are not limited to Examples of viruses, include, but are not limited to the following DNA and RNA viruses and viral families: Arbovirus, Adenoviridae, Arenaviridae, Arterivirus, Birnaviridae, Bunyaviridae, Caliciviridae, Circoviridae, Coronaviridae, Dengue, EBV, HIV, Flaviviridae, Hepadnaviridae (Hepatitis), Herpesviridae (such as, Cytomegalovirus, Herpes Simplex, Herpes Zoster), Mononegavirus (e.g., Paramyxoviridae, Morbillivirus, Rhabdoviridae), Orthomyxoviridae (e.g., Influenza A, Influenza B, and parainfluenza), Papilloma virus, Papovaviridae, Parvoviridae, Picornaviridae, Poxviridae (such as Smallpox or Vaccinia), Reoviridae (e.g., Rotavirus), Retroviridae (HTLV-I, HTLV-II, Lentivirus), and Togaviridae (e.g., Rubivirus). Viruses falling within these families can cause a variety of diseases or symptoms, including, but not limited to: arthritis, bronchiolitis, respiratory syncytial virus, encephalitis, eye infections (e.g., conjunctivitis, keratitis), chronic fatigue syndrome, hepatitis (A, B, C, E,

Chronic Active, Delta), Japanese B encephalitis, Junin, Chikungunya, Rift Valley fever, yellow fever, meningitis, opportunistic infections (e.g., AIDS), pneumonia, Burkitt's Lymphoma, chickenpox, hemorrhagic fever, Measles, Mumps, Parainfluenza, Rabies, the common cold, Polio, leukemia, Rubella, sexually transmitted diseases, skin diseases (e.g.,
5 Kaposi's, warts), and viremia. polynucleotides or polypeptides, or agonists or antagonists of the invention, can be used to treat or detect any of these symptoms or diseases. In specific embodiments, polynucleotides, polypeptides, or agonists or antagonists of the invention are used to treat: meningitis, Dengue, EBV, and/or hepatitis (e.g., hepatitis B). In an additional specific embodiment polynucleotides, polypeptides, or agonists or antagonists of the
10 invention are used to treat patients nonresponsive to one or more other commercially available hepatitis vaccines. In a further specific embodiment polynucleotides, polypeptides, or agonists or antagonists of the invention are used to treat AIDS.

Similarly, bacterial or fungal agents that can cause disease or symptoms and that can be treated or detected by a polynucleotide or polypeptide and/or agonist or antagonist of the
15 present invention include, but not limited to, include, but not limited to, the following Gram-Negative and Gram-positive bacteria and bacterial families and fungi: Actinomycetales (e.g., Corynebacterium, Mycobacterium, Norcardia), Cryptococcus neoformans, Aspergillosis, Bacillaceae (e.g., Anthrax, Clostridium), Bacteroidaceae, Blastomycosis, Bordetella, Borrelia (e.g., Borrelia burgdorferi, Brucellosis, Candidiasis, Campylobacter, Coccidioidomycosis,
20 Cryptococcosis, Dermatocycoses, E. coli (e.g., Enterotoxigenic E. coli and Enterohemorrhagic E. coli), Enterobacteriaceae (Klebsiella, Salmonella (e.g., Salmonella typhi, and Salmonella paratyphi), Serratia, Yersinia), Erysipelothrix, Helicobacter, Legionellosis, Leptospirosis, Listeria, Mycoplasmatales, Mycobacterium leprae, Vibrio cholerae, Neisseriaceae (e.g., Acinetobacter, Gonorrhea, Meningococcal), Meisseria
25 meningitidis, Pasteurellacea Infections (e.g., Actinobacillus, Heamophilus (e.g., Heamophilus influenza type B), Pasteurella), Pseudomonas, Rickettsiaceae, Chlamydiaceae, Syphilis, Shigella spp., Staphylococcal, Meningiococcal, Pneumococcal and Streptococcal (e.g., Streptococcus pneumoniae and Group B Streptococcus). These bacterial or fungal families can cause the following diseases or symptoms, including, but not limited to: bacteremia,
30 endocarditis, eye infections (conjunctivitis, tuberculosis, uveitis), gingivitis, opportunistic infections (e.g., AIDS related infections), paronychia, prosthesis-related infections, Reiter's Disease, respiratory tract infections, such as Whooping Cough or Empyema, sepsis, Lyme

Disease, Cat-Scratch Disease, Dysentery, Paratyphoid Fever, food poisoning, Typhoid, pneumonia, Gonorrhea, meningitis (e.g., meningitis types A and B), Chlamydia, Syphilis, Diphtheria, Leprosy, Paratuberculosis, Tuberculosis, Lupus, Botulism, gangrene, tetanus, impetigo, Rheumatic Fever, Scarlet Fever, sexually transmitted diseases, skin diseases (e.g., cellulitis, dermatocycoses), toxemia, urinary tract infections, wound infections. Polynucleotides or polypeptides, agonists or antagonists of the invention, can be used to treat or detect any of these symptoms or diseases. In specific embodiments, Polynucleotides, polypeptides, agonists or antagonists of the invention are used to treat: tetanus, Diphtheria, botulism, and/or meningitis type B.

Moreover, parasitic agents causing disease or symptoms that can be treated or detected by a polynucleotide or polypeptide and/or agonist or antagonist of the present invention include, but not limited to, the following families or class: Amebiasis, Babesiosis, Coccidiosis, Cryptosporidiosis, Dientamoebiasis, Dourine, Ectoparasitic, Giardiasis, Helminthiasis, Leishmaniasis, Theileriasis, Toxoplasmosis, Trypanosomiasis, and Trichomonas and Sporozoans (e.g., Plasmodium virax, Plasmodium falciparum, Plasmodium malariae and Plasmodium ovale). These parasites can cause a variety of diseases or symptoms, including, but not limited to: Scabies, Trombiculiasis, eye infections, intestinal disease (e.g., dysentery, giardiasis), liver disease, lung disease, opportunistic infections (e.g., AIDS related), malaria, pregnancy complications, and toxoplasmosis. polynucleotides or polypeptides, or agonists or antagonists of the invention, can be used to treat or detect any of these symptoms or diseases.

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention of the present invention could either be by administering an effective amount of a polypeptide to the patient, or by removing cells from the patient, supplying the cells with a polynucleotide of the present invention, and returning the engineered cells to the patient (ex vivo therapy). Moreover, the polypeptide or polynucleotide of the present invention can be used as an antigen in a vaccine to raise an immune response against infectious disease.

Regeneration

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention can be used to differentiate, proliferate, and attract cells, leading to the regeneration of tissues. (See, Science 276:59-87 (1997).) The regeneration of tissues could be used to

repair, replace, or protect tissue damaged by congenital defects, trauma (wounds, burns, incisions, or ulcers), age, disease (e.g. osteoporosis, osteoarthritis, periodontal disease, liver failure), surgery, including cosmetic plastic surgery, fibrosis, reperfusion injury, or systemic cytokine damage.

5 Tissues that could be regenerated using the present invention include organs (e.g., pancreas, liver, intestine, kidney, skin, endothelium), muscle (smooth, skeletal or cardiac), vasculature (including vascular and lymphatics), nervous, hematopoietic, and skeletal (bone, cartilage, tendon, and ligament) tissue. Preferably, regeneration occurs without or decreased scarring. Regeneration also may include angiogenesis.

10 Moreover, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, may increase regeneration of tissues difficult to heal. For example, increased tendon/ligament regeneration would quicken recovery time after damage. Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention could also be used prophylactically in an effort to avoid damage. Specific diseases that could
15 be treated include of tendinitis, carpal tunnel syndrome, and other tendon or ligament defects. A further example of tissue regeneration of non-healing wounds includes pressure ulcers, ulcers associated with vascular insufficiency, surgical, and traumatic wounds.

 Similarly, nerve and brain tissue could also be regenerated by using polynucleotides or polypeptides, as well as agonists or antagonists of the present invention, to proliferate and
20 differentiate nerve cells. Diseases that could be treated using this method include central and peripheral nervous system diseases, neuropathies, or mechanical and traumatic disorders (e.g., spinal cord disorders, head trauma, cerebrovascular disease, and stroke). Specifically, diseases associated with peripheral nerve injuries, peripheral neuropathy (e.g., resulting from chemotherapy or other medical therapies), localized neuropathies, and central nervous system
25 diseases (e.g., Alzheimer's disease, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager syndrome), could all be treated using the polynucleotides or polypeptides, as well as agonists or antagonists of the present invention.

Chem taxis

30 Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention may have chemotaxis activity. A chemotactic molecule attracts or mobilizes cells (e.g., monocytes, fibroblasts, neutrophils, T-cells, mast cells, eosinophils, epithelial and/or

endothelial cells) to a particular site in the body, such as inflammation, infection, or site of hyperproliferation. The mobilized cells can then fight off and/or heal the particular trauma or abnormality.

Polynucleotides or polypeptides, as well as agonists or antagonists of the present invention may increase chemotactic activity of particular cells. These chemotactic molecules can then be used to treat inflammation, infection, hyperproliferative disorders, or any immune system disorder by increasing the number of cells targeted to a particular location in the body. For example, chemotactic molecules can be used to treat wounds and other trauma to tissues by attracting immune cells to the injured location. Chemotactic molecules of the present invention can also attract fibroblasts, which can be used to treat wounds.

It is also contemplated that polynucleotides or polypeptides, as well as agonists or antagonists of the present invention may inhibit chemotactic activity. These molecules could also be used to treat disorders. Thus, polynucleotides or polypeptides, as well as agonists or antagonists of the present invention could be used as an inhibitor of chemotaxis.

Binding Activity

A polypeptide of the present invention may be used to screen for molecules that bind to the polypeptide or for molecules to which the polypeptide binds. The binding of the polypeptide and the molecule may activate (agonist), increase, inhibit (antagonist), or decrease activity of the polypeptide or the molecule bound. Examples of such molecules include antibodies, oligonucleotides, proteins (e.g., receptors), or small molecules.

Preferably, the molecule is closely related to the natural ligand of the polypeptide, e.g., a fragment of the ligand, or a natural substrate, a ligand, a structural or functional mimetic. (See, Coligan et al., Current Protocols in Immunology 1(2):Chapter 5 (1991).) Similarly, the molecule can be closely related to the natural receptor to which the polypeptide binds, or at least, a fragment of the receptor capable of being bound by the polypeptide (e.g., active site). In either case, the molecule can be rationally designed using known techniques.

Preferably, the screening for these molecules involves producing appropriate cells which express the polypeptide. Preferred cells include cells from mammals, yeast; *Drosophila*, or *E. coli*. Cells expressing the polypeptide (or cell membrane containing the expressed polypeptide) are then preferably contacted with a test compound potentially

containing the molecule to observe binding, stimulation, or inhibition of activity of either the polypeptide or the molecule.

The assay may simply test binding of a candidate compound to the polypeptide, wherein binding is detected by a label, or in an assay involving competition with a labeled competitor. Further, the assay may test whether the candidate compound results in a signal generated by binding to the polypeptide.

Alternatively, the assay can be carried out using cell-free preparations, polypeptide/molecule affixed to a solid support, chemical libraries, or natural product mixtures. The assay may also simply comprise the steps of mixing a candidate compound with a solution containing a polypeptide, measuring polypeptide/molecule activity or binding, and comparing the polypeptide/molecule activity or binding to a standard.

Preferably, an ELISA assay can measure polypeptide level or activity in a sample (e.g., biological sample) using a monoclonal or polyclonal antibody. The antibody can measure polypeptide level or activity by either binding, directly or indirectly, to the polypeptide or by competing with the polypeptide for a substrate.

Additionally, the receptor to which the polypeptide of the present invention binds can be identified by numerous methods known to those of skill in the art, for example, ligand panning and FACS sorting (Coligan, et al., Current Protocols in Immun., 1(2), Chapter 5, (1991)). For example, expression cloning is employed wherein polyadenylated RNA is prepared from a cell responsive to the polypeptides, for example, NIH3T3 cells which are known to contain multiple receptors for the FGF family proteins, and SC-3 cells, and a cDNA library created from this RNA is divided into pools and used to transfect COS cells or other cells that are not responsive to the polypeptides. Transfected cells which are grown on glass slides are exposed to the polypeptide of the present invention, after they have been labelled. The polypeptides can be labeled by a variety of means including iodination or inclusion of a recognition site for a site-specific protein kinase.

Following fixation and incubation, the slides are subjected to auto-radiographic analysis. Positive pools are identified and sub-pools are prepared and re-transfected using an iterative sub-pooling and re-screening process, eventually yielding a single clones that encodes the putative receptor.

As an alternative approach for receptor identification, the labeled polypeptides can be photoaffinity linked with cell membrane or extract preparations that express the receptor

molecule. Cross-linked material is resolved by PAGE analysis and exposed to X-ray film. The labeled complex containing the receptors of the polypeptides can be excised, resolved into peptide fragments, and subjected to protein microsequencing. The amino acid sequence obtained from microsequencing would be used to design a set of degenerate oligonucleotide probes to screen a cDNA library to identify the genes encoding the putative receptors.

Moreover, the techniques of gene-shuffling, motif-shuffling, exon-shuffling, and/or codon-shuffling (collectively referred to as "DNA shuffling") may be employed to modulate the activities of the polypeptide of the present invention thereby effectively generating agonists and antagonists of the polypeptide of the present invention. *See generally*, U.S. Patent Nos. 5,605,793, 5,811,238, 5,830,721, 5,834,252, and 5,837,458; and Patten, P. A., *et al.*, *Curr. Opinion Biotechnol.* 8:724-33 (1997); Harayama, S. *Trends Biotechnol.* 16(2):76-82 (1998); Hansson, L. O., *et al.*, *J. Mol. Biol.* 287:265-76 (1999); and Lorenzo, M. M. and Blasco, R. *Biotechniques* 24(2):308-13 (1998) (each of these patents and publications are hereby incorporated by reference). In one embodiment, alteration of polynucleotides and corresponding polypeptides may be achieved by DNA shuffling. DNA shuffling involves the assembly of two or more DNA segments into a desired molecule by homologous, or site-specific, recombination. In another embodiment, polynucleotides and corresponding polypeptides may be altered by being subjected to random mutagenesis by error-prone PCR, random nucleotide insertion or other methods prior to recombination. In another embodiment, one or more components, motifs, sections, parts, domains, fragments, etc., of the polypeptide of the present invention may be recombined with one or more components, motifs, sections, parts, domains, fragments, etc. of one or more heterologous molecules. In preferred embodiments, the heterologous molecules are family members. In further preferred embodiments, the heterologous molecule is a growth factor such as, for example, platelet-derived growth factor (PDGF), insulin-like growth factor (IGF-I), transforming growth factor (TGF)-alpha, epidermal growth factor (EGF), fibroblast growth factor (FGF), TGF-beta, bone morphogenetic protein (BMP)-2, BMP-4, BMP-5, BMP-6, BMP-7, activins A and B, decapentaplegic(dpp), 60A, OP-2, dorsalin, growth differentiation factors (GDFs), nodal, MIS, inhibin-alpha, TGF-beta1, TGF-beta2, TGF-beta3, TGF-beta5, and glial-derived neurotrophic factor (GDNF).

Other preferred fragments are biologically active fragments of the polypeptide of the present invention. Biologically active fragments are those exhibiting activity similar, but not

necessarily identical, to an activity of the polypeptide of the present invention. The biological activity of the fragments may include an improved desired activity, or a decreased undesirable activity.

5 Additionally, this invention provides a method of screening compounds to identify those which modulate the action of the polypeptide of the present invention. An example of such an assay comprises combining a mammalian fibroblast cell, a the polypeptide of the present invention, the compound to be screened and $^3\text{[H]}$ thymidine under cell culture conditions where the fibroblast cell would normally proliferate. A control assay may be performed in the absence of the compound to be screened and compared to the amount of
10 fibroblast proliferation in the presence of the compound to determine if the compound stimulates proliferation by determining the uptake of $^3\text{[H]}$ thymidine in each case. The amount of fibroblast cell proliferation is measured by liquid scintillation chromatography which measures the incorporation of $^3\text{[H]}$ thymidine. Both agonist and antagonist compounds may be identified by this procedure.

15 In another method, a mammalian cell or membrane preparation expressing a receptor for a polypeptide of the present invention is incubated with a labeled polypeptide of the present invention in the presence of the compound. The ability of the compound to enhance or block this interaction could then be measured. Alternatively, the response of a known second messenger system following interaction of a compound to be screened and the
20 receptor is measured and the ability of the compound to bind to the receptor and elicit a second messenger response is measured to determine if the compound is a potential agonist or antagonist. Such second messenger systems include but are not limited to, cAMP guanylate cyclase, ion channels or phosphoinositide hydrolysis.

25 All of these above assays can be used as diagnostic or prognostic markers. The molecules discovered using these assays can be used to treat disease or to bring about a particular result in a patient (e.g., blood vessel growth) by activating or inhibiting the polypeptide/molecule. Moreover, the assays can discover agents which may inhibit or enhance the production of the polypeptides of the invention from suitably manipulated cells or tissues.

30 Therefore, the invention includes a method of identifying compounds which bind to a polypeptide of the invention comprising the steps of: (a) incubating a candidate binding

compound with a polypeptide of the present invention; and (b) determining if binding has occurred. Moreover, the invention includes a method of identifying agonists/antagonists comprising the steps of: (a) incubating a candidate compound with a polypeptide of the present invention, (b) assaying a biological activity, and (b) determining if a biological activity of the polypeptide has been altered.

Targeted Delivery

In another embodiment, the invention provides a method of delivering compositions to targeted cells expressing a receptor for a polypeptide of the invention, or cells expressing a cell bound form of a polypeptide of the invention.

As discussed herein, polypeptides or antibodies of the invention may be associated with heterologous polypeptides, heterologous nucleic acids, toxins, or prodrugs via hydrophobic, hydrophilic, ionic and/or covalent interactions. In one embodiment, the invention provides a method for the specific delivery of compositions of the invention to cells by administering polypeptides of the invention (including antibodies) that are associated with heterologous polypeptides or nucleic acids. In one example, the invention provides a method for delivering a therapeutic protein into the targeted cell. In another example, the invention provides a method for delivering a single stranded nucleic acid (e.g., antisense or ribozymes) or double stranded nucleic acid (e.g., DNA that can integrate into the cell's genome or replicate episomally and that can be transcribed) into the targeted cell.

In another embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of tumor cells) by administering polypeptides of the invention (e.g., polypeptides of the invention or antibodies of the invention) in association with toxins or cytotoxic prodrugs.

By "toxin" is meant compounds that bind and activate endogenous cytotoxic effector systems, radioisotopes, holotoxins, modified toxins, catalytic subunits of toxins, or any molecules or enzymes not normally present in or on the surface of a cell that under defined conditions cause the cell's death. Toxins that may be used according to the methods of the invention include, but are not limited to, radioisotopes known in the art, compounds such as, for example, antibodies (or complement fixing containing portions thereof) that bind an inherent or induced endogenous cytotoxic effector system, thymidine kinase, endonuclease, RNase, alpha toxin, ricin, abrin, *Pseudomonas* exotoxin A, diphtheria toxin, saporin,

momordin, gelonin, pokeweed antiviral protein, alpha-sarcin and cholera toxin. By "cytotoxic prodrug" is meant a non-toxic compound that is converted by an enzyme, normally present in the cell, into a cytotoxic compound. Cytotoxic prodrugs that may be used according to the methods of the invention include, but are not limited to, glutamyl derivatives of benzoic acid mustard alkylating agent, phosphate derivatives of etoposide or
5 mitomycin C, cytosine arabinoside, daunorubisin, and phenoxyacetamide derivatives of doxorubicin.

Drug Screening

10 Further contemplated is the use of the polypeptides of the present invention, or the polynucleotides encoding these polypeptides, to screen for molecules which modify the activities of the polypeptides of the present invention. Such a method would include contacting the polypeptide of the present invention with a selected compound(s) suspected of having antagonist or agonist activity, and assaying the activity of these polypeptides
15 following binding.

This invention is particularly useful for screening therapeutic compounds by using the polypeptides of the present invention, or binding fragments thereof, in any of a variety of drug screening techniques. The polypeptide or fragment employed in such a test may be affixed to a solid support, expressed on a cell surface, free in solution, or located
20 intracellularly. One method of drug screening utilizes eukaryotic or prokaryotic host cells which are stably transformed with recombinant nucleic acids expressing the polypeptide or fragment. Drugs are screened against such transformed cells in competitive binding assays. One may measure, for example, the formulation of complexes between the agent being tested and a polypeptide of the present invention.

25 Thus, the present invention provides methods of screening for drugs or any other agents which affect activities mediated by the polypeptides of the present invention. These methods comprise contacting such an agent with a polypeptide of the present invention or a fragment thereof and assaying for the presence of a complex between the agent and the polypeptide or a fragment thereof, by methods well known in the art. In such a competitive
30 binding assay, the agents to screen are typically labeled. Following incubation, free agent is separated from that present in bound form, and the amount of free or uncomplexed label is a

measure of the ability of a particular agent to bind to the polypeptides of the present invention.

Another technique for drug screening provides high throughput screening for compounds having suitable binding affinity to the polypeptides of the present invention, and is described in great detail in European Patent Application 84/03564, published on September 13, 1984, which is incorporated herein by reference herein. Briefly stated, large numbers of different small peptide test compounds are synthesized on a solid substrate, such as plastic pins or some other surface. The peptide test compounds are reacted with polypeptides of the present invention and washed. Bound polypeptides are then detected by methods well known in the art. Purified polypeptides are coated directly onto plates for use in the aforementioned drug screening techniques. In addition, non-neutralizing antibodies may be used to capture the peptide and immobilize it on the solid support.

This invention also contemplates the use of competitive drug screening assays in which neutralizing antibodies capable of binding polypeptides of the present invention specifically compete with a test compound for binding to the polypeptides or fragments thereof. In this manner, the antibodies are used to detect the presence of any peptide which shares one or more antigenic epitopes with a polypeptide of the invention.

Antisense And Ribozyme (Antagonists)

In specific embodiments, antagonists according to the present invention are nucleic acids corresponding to the sequences contained in SEQ ID NO:X, or the complementary strand thereof, and/or to nucleotide sequences contained in a deposited cDNA clone identified in Table 1. In one embodiment, antisense sequence is generated internally, by the organism, in another embodiment, the antisense sequence is separately administered (see, for example, O'Connor, J., Neurochem. 56:560 (1991). Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988). Antisense technology can be used to control gene expression through antisense DNA or RNA, or through triple-helix formation. Antisense techniques are discussed for example, in Okano, J., Neurochem. 56:560 (1991); Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988). Triple helix formation is discussed in, for instance, Lee et al., Nucleic Acids Research 6:3073 (1979); Cooney et al., Science 241:456 (1988); and Dervan et

al., Science 251:1300 (1991). The methods are based on binding of a polynucleotide to a complementary DNA or RNA.

For example, the use of c-myc and c-myb antisense RNA constructs to inhibit the growth of the non-lymphocytic leukemia cell line HL-60 and other cell lines was previously

5 described. (Wickstrom et al. (1988); Anfossi et al. (1989)). These experiments were performed in vitro by incubating cells with the oligoribonucleotide. A similar procedure for in vivo use is described in WO 91/15580. Briefly, a pair of oligonucleotides for a given antisense RNA is produced as follows: A sequence complimentary to the first 15 bases of the open reading frame is flanked by an EcoR1 site on the 5' end and a HindIII site on the 3' end.

10 Next, the pair of oligonucleotides is heated at 90°C for one minute and then annealed in 2X ligation buffer (20mM TRIS HCl pH 7.5, 10mM MgCl₂, 10mM dithiothreitol (DTT) and 0.2 mM ATP) and then ligated to the EcoR1/Hind III site of the retroviral vector PMV7 (WO 91/15580).

For example, the 5' coding portion of a polynucleotide that encodes the polypeptide of

15 the present invention may be used to design an antisense RNA oligonucleotide of from about 10 to 40 base pairs in length. A DNA oligonucleotide is designed to be complementary to a region of the gene involved in transcription thereby preventing transcription and the production of the receptor. The antisense RNA oligonucleotide hybridizes to the mRNA in vivo and blocks translation of the mRNA molecule into receptor polypeptide.

20 In one embodiment, the antisense nucleic acid of the invention is produced intracellularly by transcription from an exogenous sequence. For example, a vector or a portion thereof, is transcribed, producing an antisense nucleic acid (RNA) of the invention. Such a vector would contain a sequence encoding the antisense nucleic acid. Such a vector can remain episomal or become chromosomally integrated, as long as it can be transcribed to

25 produce the desired antisense RNA. Such vectors can be constructed by recombinant DNA technology methods standard in the art. Vectors can be plasmid, viral, or others known in the art, used for replication and expression in vertebrate cells. Expression of the sequence encoding the polypeptide of the present invention or fragments thereof, can be by any promoter known in the art to act in vertebrate, preferably human cells. Such promoters can

30 be inducible or constitutive. Such promoters include, but are not limited to, the SV40 early promoter region (Bernoist and Chambon, Nature 29:304-310 (1981), the promoter contained in the 3' long terminal repeat of Rous sarcoma virus (Yamamoto et al., Cell 22:787-797

(1980), the herpes thymidine promoter (Wagner et al., Proc. Natl. Acad. Sci. U.S.A. 78:1441-1445 (1981), the regulatory sequences of the metallothionein gene (Brinster, et al., Nature 296:39-42 (1982)), etc.

The antisense nucleic acids of the invention comprise a sequence complementary to at least a portion of an RNA transcript of a gene of the present invention. However, absolute complementarity, although preferred, is not required. A sequence "complementary to at least a portion of an RNA," referred to herein, means a sequence having sufficient complementarity to be able to hybridize with the RNA, forming a stable duplex; in the case of double stranded antisense nucleic acids, a single strand of the duplex DNA may thus be tested, or triplex formation may be assayed. The ability to hybridize will depend on both the degree of complementarity and the length of the antisense nucleic acid. Generally, the larger the hybridizing nucleic acid, the more base mismatches with a RNA it may contain and still form a stable duplex (or triplex as the case may be). One skilled in the art can ascertain a tolerable degree of mismatch by use of standard procedures to determine the melting point of the hybridized complex.

Oligonucleotides that are complementary to the 5' end of the message, e.g., the 5' untranslated sequence up to and including the AUG initiation codon, should work most efficiently at inhibiting translation. However, sequences complementary to the 3' untranslated sequences of mRNAs have been shown to be effective at inhibiting translation of mRNAs as well. See generally, Wagner, R., 1994, Nature 372:333-335. Thus, oligonucleotides complementary to either the 5'- or 3'- non- translated, non-coding regions of polynucleotide sequences described herein could be used in an antisense approach to inhibit translation of endogenous mRNA. Oligonucleotides complementary to the 5' untranslated region of the mRNA should include the complement of the AUG start codon. Antisense oligonucleotides complementary to mRNA coding regions are less efficient inhibitors of translation but could be used in accordance with the invention. Whether designed to hybridize to the 5', 3' or coding region of mRNA of the present invention, antisense nucleic acids should be at least six nucleotides in length, and are preferably oligonucleotides ranging from 6 to about 50 nucleotides in length. In specific aspects the oligonucleotide is at least 10 nucleotides, at least 17 nucleotides, at least 25 nucleotides or at least 50 nucleotides.

The polynucleotides of the invention can be DNA or RNA or chimeric mixtures or derivatives or modified versions thereof, single-stranded or double-stranded. The

oligonucleotide can be modified at the base moiety, sugar moiety, or phosphate backbone, for example, to improve stability of the molecule, hybridization, etc. The oligonucleotide may include other appended groups such as peptides (e.g., for targeting host cell receptors in vivo), or agents facilitating transport across the cell membrane (see, e.g., Letsinger et al., 1989, Proc. Natl. Acad. Sci. U.S.A. 86:6553-6556; Lemaitre et al., 1987, Proc. Natl. Acad. Sci. 84:648-652; PCT Publication No. WO88/09810, published December 15, 1988) or the blood-brain barrier (see, e.g., PCT Publication No. WO89/10134, published April 25, 1988), hybridization-triggered cleavage agents. (See, e.g., Krol et al., 1988, BioTechniques 6:958-976) or intercalating agents. (See, e.g., Zon, 1988, Pharm. Res. 5:539-549). To this end, the oligonucleotide may be conjugated to another molecule, e.g., a peptide, hybridization triggered cross-linking agent, transport agent, hybridization-triggered cleavage agent, etc.

The antisense oligonucleotide may comprise at least one modified base moiety which is selected from the group including, but not limited to, 5-fluorouracil, 5-bromouracil, 5-chlorouracil, 5-iodouracil, hypoxanthine, xantine, 4-acetylcytosine, 5-(carboxyhydroxymethyl) uracil, 5-carboxymethylaminomethyl-2-thiouridine, 5-carboxymethylaminomethyluracil, dihydrouracil, beta-D-galactosylqueosine, inosine, N6-isopentenyladenine, 1-methylguanine, 1-methylinosine, 2,2-dimethylguanine, 2-methyladenine, 2-methylguanine, 3-methylcytosine, 5-methylcytosine, N6-adenine, 7-methylguanine, 5-methylaminomethyluracil, 5-methoxyaminomethyl-2-thiouracil, beta-D-mannosylqueosine, 5'-methoxycarboxymethyluracil, 5-methoxyuracil, 2-methylthio-N6-isopentenyladenine, uracil-5-oxyacetic acid (v), wybutoxosine, pseudouracil, queosine, 2-thiocytosine, 5-methyl-2-thiouracil, 2-thiouracil, 4-thiouracil, 5-methyluracil, uracil-5-oxyacetic acid methylester, uracil-5-oxyacetic acid (v), 5-methyl-2-thiouracil, 3-(3-amino-3-N-2-carboxypropyl) uracil, (acp3)w, and 2,6-diaminopurine.

The antisense oligonucleotide may also comprise at least one modified sugar moiety selected from the group including, but not limited to, arabinose, 2-fluoroarabinose, xylulose, and hexose.

In yet another embodiment, the antisense oligonucleotide comprises at least one modified phosphate backbone selected from the group including, but not limited to, a phosphorothioate, a phosphorodithioate, a phosphoramidothioate, a phosphoramidate, a phosphordiamidate, a methylphosphonate, an alkyl phosphotriester, and a formacetal or analog thereof.

In yet another embodiment, the antisense oligonucleotide is an a-anomeric oligonucleotide. An a-anomeric oligonucleotide forms specific double-stranded hybrids with complementary RNA in which, contrary to the usual b-units, the strands run parallel to each other (Gautier et al., 1987, Nucl. Acids Res. 15:6625-6641). The oligonucleotide is a 2'-O-methylribonucleotide (Inoue et al., 1987, Nucl. Acids Res. 15:6131-6148), or a chimeric RNA-DNA analogue (Inoue et al., 1987, FEBS Lett. 215:327-330).

Polynucleotides of the invention may be synthesized by standard methods known in the art, e.g. by use of an automated DNA synthesizer (such as are commercially available from Biosearch, Applied Biosystems, etc.). As examples, phosphorothioate oligonucleotides may be synthesized by the method of Stein et al. (1988, Nucl. Acids Res. 16:3209), methylphosphonate oligonucleotides can be prepared by use of controlled pore glass polymer supports (Sarin et al., 1988, Proc. Natl. Acad. Sci. U.S.A. 85:7448-7451), etc.

While antisense nucleotides complementary to the coding region sequence could be used, those complementary to the transcribed untranslated region are most preferred.

Potential antagonists according to the invention also include catalytic RNA, or a ribozyme (See, e.g., PCT International Publication WO 90/11364, published October 4, 1990; Sarver et al, Science 247:1222-1225 (1990). While ribozymes that cleave mRNA at site specific recognition sequences can be used to destroy mRNAs, the use of hammerhead ribozymes is preferred. Hammerhead ribozymes cleave mRNAs at locations dictated by flanking regions that form complementary base pairs with the target mRNA. The sole requirement is that the target mRNA have the following sequence of two bases: 5'-UG-3'. The construction and production of hammerhead ribozymes is well known in the art and is described more fully in Haseloff and Gerlach, Nature 334:585-591 (1988). There are numerous potential hammerhead ribozyme cleavage sites within the nucleotide sequence of SEQ ID NO:X. Preferably, the ribozyme is engineered so that the cleavage recognition site is located near the 5' end of the mRNA; i.e., to increase efficiency and minimize the intracellular accumulation of non-functional mRNA transcripts.

As in the antisense approach, the ribozymes of the invention can be composed of modified oligonucleotides (e.g. for improved stability, targeting, etc.) and should be delivered to cells which express in vivo. DNA constructs encoding the ribozyme may be introduced into the cell in the same manner as described above for the introduction of antisense encoding DNA. A preferred method of delivery involves using a DNA construct "encoding" the

ribozyme under the control of a strong constitutive promoter, such as, for example, pol III or pol II promoter, so that transfected cells will produce sufficient quantities of the ribozyme to destroy endogenous messages and inhibit translation. Since ribozymes unlike antisense molecules, are catalytic, a lower intracellular concentration is required for efficiency.

5 Antagonist/agonist compounds may be employed to inhibit the cell growth and proliferation effects of the polypeptides of the present invention on neoplastic cells and tissues, i.e. stimulation of angiogenesis of tumors, and, therefore, retard or prevent abnormal cellular growth and proliferation, for example, in tumor formation or growth, particularly in prostate tumor formation or growth.

10 The antagonist/agonist may also be employed to prevent hyper-vascular diseases, and prevent the proliferation of epithelial lens cells after extracapsular cataract surgery. Prevention of the mitogenic activity of the polypeptides of the present invention may also be desirable in cases such as restenosis after balloon angioplasty.

The antagonist/agonist may also be employed to prevent the growth of scar tissue
15 during wound healing.

The antagonist/agonist may also be employed to treat the diseases described herein.

Thus, the invention provides a method of treating disorders or diseases, including but not limited to the disorders or diseases listed throughout this application, associated with overexpression of a polynucleotide of the present invention by administering to a patient (a)
20 an antisense molecule directed to the polynucleotide of the present invention, and/or (b) a ribozyme directed to the polynucleotide of the present invention.

Other Activities

A polypeptide, polynucleotide, agonist, or antagonist of the present invention, as a
25 result of the ability to stimulate vascular endothelial cell growth, may be employed in treatment for stimulating re-vascularization of ischemic tissues due to various disease conditions such as thrombosis, arteriosclerosis, and other cardiovascular conditions. The polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be employed to stimulate angiogenesis and limb regeneration, as discussed above.

30 A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be employed for treating wounds due to injuries, burns, post-operative tissue repair, and ulcers since they are mitogenic to various cells of different origins, such as fibroblast cells

and skeletal muscle cells, and therefore, facilitate the repair or replacement of damaged or diseased tissue.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be employed stimulate neuronal growth and to treat and prevent neuronal damage which occurs in certain neuronal disorders or neuro-degenerative conditions such as Alzheimer's disease, Parkinson's disease, and AIDS-related complex. A polypeptide, polynucleotide, agonist, or antagonist of the present invention may have the ability to stimulate chondrocyte growth, therefore, they may be employed to enhance bone and periodontal regeneration and aid in tissue transplants or bone grafts.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may be also be employed to prevent skin aging due to sunburn by stimulating keratinocyte growth.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be employed for preventing hair loss, since FGF family members activate hair-forming cells and promotes melanocyte growth. Along the same lines, a polypeptide, polynucleotide, agonist, or antagonist of the present invention may be employed to stimulate growth and differentiation of hematopoietic cells and bone marrow cells when used in combination with other cytokines.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be employed to maintain organs before transplantation or for supporting cell culture of primary tissues. A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be employed for inducing tissue of mesodermal origin to differentiate in early embryos.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also increase or decrease the differentiation or proliferation of embryonic stem cells, besides, as discussed above, hematopoietic lineage.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be used to modulate mammalian characteristics, such as body height, weight, hair color, eye color, skin, percentage of adipose tissue, pigmentation, size, and shape (e.g., cosmetic surgery). Similarly, a polypeptide, polynucleotide, agonist, or antagonist of the present invention may be used to modulate mammalian metabolism affecting catabolism, anabolism, processing, utilization, and storage of energy.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may be used to change a mammal's mental state or physical state by influencing biorhythms, cardiac rhythms, depression (including depressive disorders), tendency for violence, tolerance for pain, reproductive capabilities (preferably by Activin or Inhibin-like activity),
5 hormonal or endocrine levels, appetite, libido, memory, stress, or other cognitive qualities.

A polypeptide, polynucleotide, agonist, or antagonist of the present invention may also be used as a food additive or preservative, such as to increase or decrease storage capabilities, fat content, lipid, protein, carbohydrate, vitamins, minerals, cofactors or other nutritional components.

10 The above-recited applications have uses in a wide variety of hosts. Such hosts include, but are not limited to, human, murine, rabbit, goat, guinea pig, camel, horse, mouse, rat, hamster, pig, micro-pig, chicken, goat, cow, sheep, dog, cat, non-human primate, and human. In specific embodiments, the host is a mouse, rabbit, goat, guinea pig, chicken, rat, hamster, pig, sheep, dog or cat. In preferred embodiments, the host is a mammal. In most
15 preferred embodiments, the host is a human.

Other Preferred Embodiments

Other preferred embodiments of the claimed invention include an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to a sequence
20 of at least about 50 contiguous nucleotides in the nucleotide sequence of SEQ ID NO:X wherein X is any integer as defined in Table 1.

Also preferred is a nucleic acid molecule wherein said sequence of contiguous nucleotides is included in the nucleotide sequence of SEQ ID NO:X in the range of positions beginning with the nucleotide at about the position of the 5' Nucleotide of the Clone
25 Sequence and ending with the nucleotide at about the position of the 3' Nucleotide of the Clone Sequence as defined for SEQ ID NO:X in Table 1.

Also preferred is a nucleic acid molecule wherein said sequence of contiguous nucleotides is included in the nucleotide sequence of SEQ ID NO:X in the range of positions beginning with the nucleotide at about the position of the 5' Nucleotide of the Start Codon
30 and ending with the nucleotide at about the position of the 3' Nucleotide of the Clone Sequence as defined for SEQ ID NO:X in Table 1, or the complementary strand thereto.

Also preferred is an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to a sequence of at least about 150 contiguous nucleotides in the nucleotide sequence of SEQ ID NO:X or the complementary strand thereto.

Further preferred is an isolated nucleic acid molecule comprising a nucleotide
5 sequence which is at least 95% identical to a sequence of at least about 500 contiguous nucleotides in the nucleotide sequence of SEQ ID NO:X, or the complementary strand thereto.

A further preferred embodiment is a nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to the nucleotide sequence of SEQ ID NO:X
10 beginning with the nucleotide at about the position of the 5' Nucleotide of the First Amino Acid and ending with the nucleotide at about the position of the 3' Nucleotide of the Clone Sequence as defined for SEQ ID NO:X in Table 1, or the complementary strand thereto.

A further preferred embodiment is an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to the complete nucleotide sequence of
15 SEQ ID NO:X, or the complementary strand thereto.

Also preferred is an isolated nucleic acid molecule which hybridizes under stringent hybridization conditions to a nucleic acid molecule comprising a nucleotide sequence of SEQ ID NO:X or the complementary strand thereto and/or cDNA contained in ATCC Deposit No.:Z, wherein said nucleic acid molecule which hybridizes does not hybridize under
20 stringent hybridization conditions to a nucleic acid molecule having a nucleotide sequence consisting of only A residues or of only T residues.

Also preferred is a composition of matter comprising a DNA molecule which comprises a human cDNA clone identified by a cDNA Clone Identifier in Table 1, which DNA molecule is contained in the material deposited with the American Type Culture
25 Collection and given ATCC Deposit No.:Z.

Also preferred is a composition of matter comprising a DNA molecule which comprises the cDNA contained in ATCC Deposit No.:Z.

Also preferred is an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to a sequence of at least 50 contiguous nucleotides of the
30 cDNA sequence contained in ATCC Deposit No.:Z.

Also preferred is an isolated nucleic acid molecule, wherein said sequence of at least 50 contiguous nucleotides is included in the nucleotide sequence of an open reading frame sequence encoded by cDNA sequence contained in ATCC Deposit No.:Z.

Also preferred is an isolated nucleic acid molecule comprising a nucleotide sequence
5 which is at least 95% identical to sequence of at least 150 contiguous nucleotides in the nucleotide sequence encoded by cDNA sequence contained in ATCC Deposit No.:Z.

A further preferred embodiment is an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to sequence of at least 500 contiguous nucleotides in the nucleotide sequence encoded by cDNA sequence contained in ATCC
10 Deposit No.:Z.

A further preferred embodiment is an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to the complete nucleotide sequence encoded by said human cDNA clone.

A further preferred embodiment is a method for detecting in a biological sample a
15 nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to a sequence of at least 50 contiguous nucleotides in a sequence selected from the group consisting of: a nucleotide sequence of SEQ ID NO:X wherein X is any integer as defined in Table 1 or the complementary strand thereto; and a nucleotide sequence encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC
20 Deposit No.:Z; which method comprises a step of comparing a nucleotide sequence of at least one nucleic acid molecule in said sample with a sequence selected from said group and determining whether the sequence of said nucleic acid molecule in said sample is at least 95% identical to said selected sequence.

Also preferred is the above method wherein said step of comparing sequences
25 comprises determining the extent of nucleic acid hybridization between nucleic acid molecules in said sample and a nucleic acid molecule comprising said sequence selected from said group. Similarly, also preferred is the above method wherein said step of comparing sequences is performed by comparing the nucleotide sequence determined from a nucleic acid molecule in said sample with said sequence selected from said group. The nucleic acid
30 molecules can comprise DNA molecules or RNA molecules.

A further preferred embodiment is a method for identifying the species, tissue or cell type of a biological sample which method comprises a step of detecting nucleic acid

molecules in said sample, if any, comprising a nucleotide sequence that is at least 95% identical to a sequence of at least 50 contiguous nucleotides in a sequence selected from the group consisting of: a nucleotide sequence of SEQ ID NO:X wherein X is any integer as defined in Table 1 or the complement thereto; and a nucleotide sequence encoded by a human
5 cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

The method for identifying the species, tissue or cell type of a biological sample can comprise a step of detecting nucleic acid molecules comprising a nucleotide sequence in a panel of at least two nucleotide sequences, wherein at least one sequence in said panel is at
10 least 95% identical to a sequence of at least 50 contiguous nucleotides in a sequence selected from said group.

Also preferred is a method for diagnosing in a subject a pathological condition associated with abnormal structure or expression of a gene encoding a protein identified in Table 1, wherein the method comprises a step of detecting in a biological sample obtained
15 from said subject nucleic acid molecules, if any, comprising a nucleotide sequence that is at least 95% identical to a sequence of at least 50 contiguous nucleotides in a sequence selected from the group consisting of: a nucleotide sequence of SEQ ID NO:X wherein X is any integer as defined in Table 1 or the complementary strand thereto; and a nucleotide sequence encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and
20 contained in ATCC Deposit No.:Z.

The method for diagnosing a pathological condition can comprise a step of detecting nucleic acid molecules comprising a nucleotide sequence in a panel of at least two nucleotide sequences, wherein at least one sequence in said panel is at least 95% identical to a sequence
of at least 50 contiguous nucleotides in a sequence selected from said group.

25 Also preferred is a composition of matter comprising isolated nucleic acid molecules wherein the nucleotide sequences of said nucleic acid molecules comprise a panel of at least two nucleotide sequences, wherein at least one sequence in said panel is at least 95% identical to a sequence of at least 50 contiguous nucleotides in a sequence selected from the group consisting of: a nucleotide sequence of SEQ ID NO:X wherein X is any integer as
30 defined in Table 1 or the complementary strand thereto; and a nucleotide sequence encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in

ATCC Deposit No.:Z. The nucleic acid molecules can comprise DNA molecules or RNA molecules.

Also preferred is a composition of matter comprising isolated nucleic acid molecules wherein the nucleotide sequences of said nucleic acid molecules comprise a DNA microarray or "chip" of at least 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 40, 50, 100, 150, 200, 250, 300, 500, 1000, 2000, 3000 or 4000 nucleotide sequences, wherein at least one sequence in said DNA microarray or "chip" is at least 95% identical to a sequence of at least 50 contiguous nucleotides in a sequence selected from the group consisting of: a nucleotide sequence of SEQ ID NO:X wherein X is any integer as defined in Table 1; and a nucleotide sequence encoded by a human cDNA clone identified in Table 1 which was deposited in ATCC Deposit No.:Z. The nucleic acid molecules can comprise DNA molecules or RNA molecules.

Also preferred is an isolated polypeptide comprising an amino acid sequence at least 90% identical to a sequence of at least about 10 contiguous amino acids in the amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1, and/or a polypeptide encoded by SEQ ID NO:X.

Also preferred is an isolated polypeptide comprising an amino acid sequence at least 95% identical to a sequence of at least about 30 contiguous amino acids in the amino acid sequence of SEQ ID NO:Y and/or a polypeptide encoded by SEQ ID NO:X.

Further preferred is an isolated polypeptide comprising an amino acid sequence at least 95% identical to a sequence of at least about 100 contiguous amino acids in the amino acid sequence of SEQ ID NO:Y and/or a polypeptide encoded by SEQ ID NO:X.

Further preferred is an isolated polypeptide comprising an amino acid sequence at least 95% identical to the complete amino acid sequence of SEQ ID NO:Y and/or a polypeptide encoded by SEQ ID NO:X.

Further preferred is an isolated polypeptide comprising an amino acid sequence at least 90% identical to a sequence of at least about 10 contiguous amino acids in the complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is a polypeptide wherein said sequence of contiguous amino acids is included in the amino acid sequence of a portion of the protein encoded by a human cDNA

clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is an isolated polypeptide comprising an amino acid sequence at least 95% identical to a sequence of at least about 30 contiguous amino acids in the amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is an isolated polypeptide comprising an amino acid sequence at least 95% identical to a sequence of at least about 100 contiguous amino acids in the amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is an isolated polypeptide comprising an amino acid sequence at least 95% identical to the amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Further preferred is an isolated antibody which binds specifically to a polypeptide comprising an amino acid sequence that is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; an amino acid sequence encoded by SEQ ID NO:X and a complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Further preferred is a method for detecting in a biological sample a polypeptide comprising an amino acid sequence which is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; an amino acid sequence encoded by SEQ ID NO:X, and a complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z; which method comprises a step of comparing an amino acid sequence of at least one polypeptide molecule in said sample with a sequence selected from said group and determining whether the sequence of said polypeptide molecule in said sample is at least 90% identical to said sequence of at least 10 contiguous amino acids.

Also preferred is the above method wherein said step of comparing an amino acid sequence of at least one polypeptide molecule in said sample with a sequence selected from

said group comprises determining the extent of specific binding of polypeptides in said sample to an antibody which binds specifically to a polypeptide comprising an amino acid sequence that is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; an amino acid sequence encoded by SEQ ID NO:X, and a complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is the above method wherein said step of comparing sequences is performed by comparing the amino acid sequence determined from a polypeptide molecule in said sample with said sequence selected from said group.

Also preferred is a method for identifying the species, tissue or cell type of a biological sample which method comprises a step of detecting polypeptide molecules in said sample, if any, comprising an amino acid sequence that is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; and amino acid sequence encoded by SEQ ID NO:X, and a complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is the above method for identifying the species, tissue or cell type of a biological sample, which method comprises a step of detecting polypeptide molecules comprising an amino acid sequence in a panel of at least two amino acid sequences, wherein at least one sequence in said panel is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the above group.

Also preferred is a method for diagnosing in a subject a pathological condition associated with abnormal structure or expression of a gene encoding a protein identified in Table 1, which method comprises a step of detecting in a biological sample obtained from said subject polypeptide molecules comprising an amino acid sequence in a panel of at least two amino acid sequences, wherein at least one sequence in said panel is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; and a complete amino acid sequence of a protein encoded by a human

cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

In any of these methods, the step of detecting said polypeptide molecules includes using an antibody.

5 Also preferred is an isolated nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to a nucleotide sequence encoding a polypeptide wherein said polypeptide comprises an amino acid sequence that is at least 90% identical to a sequence of at least 10 contiguous amino acids in a sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; and a
10 complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

Also preferred is an isolated nucleic acid molecule, wherein said nucleotide sequence encoding a polypeptide has been optimized for expression of said polypeptide in a prokaryotic host.

15 Also preferred is an isolated nucleic acid molecule, wherein said polypeptide comprises an amino acid sequence selected from the group consisting of: an amino acid sequence of SEQ ID NO:Y wherein Y is any integer as defined in Table 1; and a complete amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z.

20 Further preferred is a method of making a recombinant vector comprising inserting any of the above isolated nucleic acid molecule into a vector. Also preferred is the recombinant vector produced by this method. Also preferred is a method of making a recombinant host cell comprising introducing the vector into a host cell, as well as the recombinant host cell produced by this method.

25 Also preferred is a method of making an isolated polypeptide comprising culturing this recombinant host cell under conditions such that said polypeptide is expressed and recovering said polypeptide. Also preferred is this method of making an isolated polypeptide, wherein said recombinant host cell is a eukaryotic cell and said polypeptide is a human protein comprising an amino acid sequence selected from the group consisting of: an
30 amino acid sequence of SEQ ID NO:Y beginning with the residue at the position of the First Amino Acid of the Protein of SEQ ID NO:Y wherein Y is an integer set forth in Table 1 and said position of the First Amino Acid of the Protein of SEQ ID NO:Y is defined in Table 1;

and an amino acid sequence of a protein encoded by a human cDNA clone identified by a cDNA Clone Identifier in Table 1 and contained in ATCC Deposit No.:Z. The isolated polypeptide produced by this method is also preferred.

Also preferred is a method of treatment of an individual in need of an increased level of a protein activity, which method comprises administering to such an individual a Therapeutic comprising an amount of an isolated polypeptide, polynucleotide, immunogenic fragment or analogue thereof, binding agent, antibody, or antigen binding fragment of the claimed invention effective to increase the level of said protein activity in said individual.

Also preferred is a method of treatment of an individual in need of a decreased level of a protein activity, which method comprised administering to such an individual a Therapeutic comprising an amount of an isolated polypeptide, polynucleotide, immunogenic fragment or analogue thereof, binding agent, antibody, or antigen binding fragment of the claimed invention effective to decrease the level of said protein activity in said individual.

Also preferred is a method of treatment of an individual in need of specific delivery of toxic compositions to diseased cells (e.g., cancers or tumors, including but not limited to prostate cancer cells, leukemias or lymphomas), which method comprises administering to such an individual a Therapeutic comprising an amount of an isolated polypeptide of the invention, including but not limited to a binding agent, or antibody of the claimed invention that are associated with toxin or cytotoxic prodrugs.

In specific embodiments of the invention, for each "Contig ID" listed in the fourth column of Table 2, preferably excluded are one or more polynucleotides comprising, or alternatively consisting of, a nucleotide sequence referenced in the fifth column of Table 2 and described by the general formula of a-b, whereas a and b are uniquely determined for the corresponding SEQ ID NO:X referred to in column 3 of Table 2. Further specific embodiments are directed to polynucleotide sequences excluding one, two, three, four, or more of the specific polynucleotide sequences referred to in the fifth column of Table 2. In no way is this listing meant to encompass all of the sequences which may be excluded by the general formula, it is just a representative example. All references available through these accessions are hereby incorporated by reference in their entirety.

TABLE II

Gene N .	cDNA Cl ne ID	NT SEQ	Contig ID	Public Accession Num b rs
----------	------------------	-----------	-----------	---------------------------

		ID NO: X		
1	HHEPE84	11	859463	None
1	HHEPE84	36	840809	H12690, H99220, H99814, N40464, N48503, W49785, AA190704, AA191524, AA463219, AA430306, AA805957, AA776841, AA812641, AA843424, AA904606
2	HTXFS09	12	866494	None
3	HSPAF01	13	862062	None
3	HSPAF01	37	840747	T50792, T53826, T53958, T60312, T60333, T61068, T61104, T61132, T61171, T61626, T64577, T67689, T68882, T69443, T73181, T74749, T84061, T86099, T97646, R72285, R72801, R73546, H21515, H21514, H27643, H28077, H30180, H42471, H42674, H42726, H43452, H44632, H46408, R86068, R97964, R97965, R99755, H60221, H61128, H78168, H78169, H79387, N73598, W02564, W77765, W78004, W79463, W94583, W95318, AA075014, AA143598
4	HWHHB29, HSRAA55	14	1031953	R27201, R27200, R53918, R53919, H10706, H92023, H92022, N48159, N51224, N53305, N59016, W63643, AA043242, AA043571, AA081648, AA174144, AA176689, AA424157, AA279572, AA601516, AA604338, AA578387, AA657498, AA732358, AA737454, AA746000, AA760751, AA806853, AA877968, AA934492, AA974173, D82793, N56274, C01847, C05245, AA094783, AA643624, AA654027, Z19471, AA852141, AA852140, AA884822, AA916315, AI041376, Z39202, T16750, AA952949, T27278, AA693381, AI241213, AI310741, AI355499, AI439466, AI151477, AI263704, AI538941, AI591369, AI521934, AI769711, AI858795, AI955847, AW026715, AW088807, AW276257, AW303919, AW305223, AW439725
4	HWHHB29, HSRAA55	38	1042279	None
5	HLWBY67	15	859469	None
5	HLWBY67	39	829086	None
5	HLWBY67	40	829087	R25382, R28405, R46543, R74048, R74141, R80763, R80762, H00272, H00273, H24568, H24590, H27776, H27844, H39888, H42200, H42269, N93392, W71996, W76510, AA130428, AA130429, AA195178, AA463354, AA463366, AA463861, AA463870, AA422151, AA480665, AA507859, AA532812, AA552205, AA583338, AA588102, AA568654, AA627351, AA635438, AA640268, AA579639, AA808628, AA812319, AA826714, AA838506, AA837856, AA858248, AI075002
6	HPWBU56, HEBFL69	16	886908	AA226171, AA484674, AA502194, AA525269, AA532647, AA662702, AA662733, AI198311, AI198986, AI199681, AI291430
6	HPWBU56,	41	866479	None

	HEBFL69			
6	HPWBU56, HEBFL69	42	828649	AA226171, AA484674, AA502194, AA525269, AA532647, AA662702, AA662733
7	HCFBC32	17	862004	None
7	HCFBC32	43	841198	H40835, H71461, H71590, H71461, W93450, W95880, W95916, AA009722, AA009418, AA132972, AA143286, AA149432, AA151515, AA166705, AA166775, AA213925, AA283826, AA284064, AA213627, AA465150, AA465258, AA505421, AA534052, AA552763, AA555201, AA587097, AA640568, AA576127, AA657429, AA659320, AA661655, AA745066, AA826877, AA857750, AA888269, AA913282, AA987858, AI083826, W22226, W22528, AA143230, AA652536, AA479114, AA479264, AA707128, AA719823, AA774855, AA775593, AI014998, D20958, AA694580, AA700694, AA700715
8	HSLJW05	18	866210	None
8	HSLJW05	44	840739	T80169, T80449, R24660, R25163, R39019, R39369, R62525, R82222, AA214516, AA280756, AA528458, AA636097, AA878377, AA969711, AA995304, C16437, AI084502, AI086537, D20028
9	HJAAT54	19	877578	None
9	HJAAT54	45	637706	None
10	HMEFI17, HMEFK17	20	889121	None
11	hdpqe64, HCEES14, HDPQE64	21	873791	T48922, T52896, T54452, T48923, T52897, T54514, T82450, R22428, R22478, H67812, H68084, H87606, H87605, N22404, N30588, N45519, N47617, N57233, N57232, N70370, W19722, N89978, AA017666, AA045567, AA057053, AA427619, AA558095, AI088764, W22860, AI097118, AA018850, C18158, C18797, C17709, AA437353, AA720001, AA720020, AA897572, AI033301, AI078459, F09978, AI337140, AI361611, AI365337, AI431831, AI494034, AI565998, AI123818, AI126025, AI189726, AI189724, AI656384, AI219209, AI611185, AI669975
11	hdpqe64, HCEES14, HDPQE64	46	841067	None
11	hdpqe64, HCEES14, HDPQE64	47	829626	T48922, T52896, T54452, T48923, T52897, T54514, T82450, R22428, R22478, H67812, H68084, H87606, H87605, N22404, N30588, N45519, N47617, N57233, N57232, N70370, W19722, N89978, AA017666, AA045567, AA057053, AA427619
12	HISAN67	22	866451	None
12	HISAN67	48	829296	H47409, H77806, H87648, H89639, H90940, H95471, N99357, W02522
13	HMWJD58, HCRBF72	23	873773	R02577, R02696, R54743, W47056, W77999, W79458, AA076651, AA079559, AA135359, AA137236, AA172174, AA808954, AA904052, AA974194, C06498, AA293812, AA402460, F20535, AA680169, AA993790, AI085220,

				D20372, AI318369, AI365185, AI346937, AI202349, AI499014, AI146448, AI185138, AI652019, AI219673, AI636880, AI638001, AI632130
14	HMQAI69	24	877830	None
14	HMQAI69	49	829009	H17108, H26527, H95758, N62859, N77765, AA425013, F16426, AA582507, AA789136, AA830141, AA927512, AA642790
15	HRAAA23	25	872460	None
15	HRAAA23	50	828542	AA928914
16	HOGCK09	26	877831	None
16	HOGCK09	51	809101	None
17	HPRAT22	27	866475	None
17	HPRAT22	52	828625	AA946965
18	HTXDT74	28	872573	None
18	HTXDT74	53	840625	None
19	HDPRJ46	29	841052	R10233, R10739, R25387, R28410, R35893, R35894, R82187, R82235, H01419, H01420, H02059, H04908, H05883, H47909, H47910, R86249, R86250, R97085, R97132, R98809, H48386, H78551, H79038, H82693, H86846, N29331, N29336, N30236, N57023, N57029, N57128, N62670, N73914, N80444, W03978, W31268, W37581, AA011095, AA011096, AA035648, AA035752, AA514315, AA533001, AA535101, AA573451, AA865791, AA873797, AA873246, AA878907, AA878660, AA995194, AA723979, AI057254, AI057372, AI057376, AI076430, AI076531, F06002
19	HDPRJ46	54	859464	None
20	HEQAN39	30	859468	None
20	HEQAN39	55	840923	R19882, R45106, R45106, H06732, H11060, H11146, H39503, R92995, H72442, H95661, N21600, N22604, N28833, N30271, N35792, N41993, N49950, N49960, N50639, N50720, N52731, N67962, N73183, N74910, W23515, W67163, W67164, AA121018, AA126466, AA226162, AA504324, AA600768, AA614188, AA627363, AA824469, AA917360, AA932053, AA999988, N55398, C14171, AA219351, AA410507, AA453255, AA453356, AA854303, AA885069, AA910469, AA971667, AA985055, AI017619, AI051010, T17235, Z40061, Z46061, AA699500
21	HFKFN13	31	866445	None
21	HFKFN13	56	840870	T50014, T50076, T77457, R10123, R10531, R11582, R13738, R19301, R20773, R37148, R83520, H52002, H52120, N24189, N31678, N57536, N70170, N71169, N98366, W05521, W21341, W58727, W68327, W74313, W79247, W79304, W79299, AA023011, AA022477, AA029469, AA029536, AA034257, AA063534, AA173286, AA173210, AA252660, AA252659, AA465008, AA491386, AA504775, L44428, AA828515, AA847841, AA922310, AA938769, AA975821, AA641879, AA641923, AA449413,

				AA450379, AA455288, AA626495, AA626488, AA778184, AA984253, AI015506, AI076206, AI122921, D20333, Z41249, Z45530, Z45553
22	HPFDD04	32	866474	None
22	HPFDD04	57	828768	R09227, R09339, AA491988, AA507505
23	HPWDE02	33	877829	None
23	HPWDE02	58	828544	AA216357, AA226583, AA229229, AA229263, AA229408, AA229424, AA228731, AA228669, AA630840, AA639902, AA579452, AA688095
24	HPIBS12	34	866105	None
24	HPIBS12	59	828687	None
25	HPIAD06	35	872588	None

Having generally described the invention, the same will be more readily understood by reference to the following examples, which are provided by way of illustration and are not intended as limiting.

*Examples**Example 1: Isolation of a Selected cDNA Clone From the Deposited Sample*

5 Each cDNA clone in a cited ATCC deposit is contained in a plasmid vector. Table 1 identifies the vectors used to construct the cDNA library from which each clone was isolated. In many cases, the vector used to construct the library is a phage vector from which a plasmid has been excised. The information immediately below correlates the related plasmid for each phage vector used in constructing the cDNA library. For example, where a particular clone is
 10 identified in Table 1 as being isolated in the vector "Lambda Zap," the corresponding deposited clone is in "pBluescript."

	<u>Vector Used to Construct Library</u>	<u>Corresponding Deposited Plasmid</u>
	Lambda Zap	pBluescript (pBS)
	Uni-Zap XR	pBluescript (pBS)
15	Zap Express	pBK
	lafmid BA	plafmid BA
	pSport1	pSport1
	pCMVSPORT 2.0	pCMVSPORT 2.0
	pCMVSPORT 3.0	pCMVSPORT 3.0
20	pCR [®] 2.1	pCR [®] 2.1

Vectors Lambda Zap (U.S. Patent Nos. 5,128,256 and 5,286,636), Uni-Zap XR (U.S. Patent Nos. 5,128, 256 and 5,286,636), Zap Express (U.S. Patent Nos. 5,128,256 and 5,286,636), pBluescript (pBS) (Short, J. M. et al., Nucleic Acids Res. 16:7583-7600 (1988); Alting-Mees, M. A. and Short, J. M., Nucleic Acids Res. 17:9494 (1989)) and pBK (Alting-Mees, M. A. et al., Strategies 5:58-61 (1992)) are commercially available from Stratagene
 25 Cloning Systems, Inc., 11011 N. Torrey Pines Road, La Jolla, CA, 92037. pBS contains an ampicillin resistance gene and pBK contains a neomycin resistance gene. Both can be transformed into E. coli strain XL-1 Blue, also available from Stratagene. pBS comes in 4 forms SK+, SK-, KS+ and KS. The S and K refers to the orientation of the polylinker to the
 30 T7 and T3 primer sequences which flank the polylinker region ("S" is for SacI and "K" is for KpnI which are the first sites on each respective end of the linker). "+" or "-" refer to the

orientation of the fl origin of replication ("ori"), such that in one orientation, single stranded rescue initiated from the fl ori generates sense strand DNA and in the other, antisense.

Vectors pSport1, pCMVSPORT 2.0 and pCMVSPORT 3.0, were obtained from Life Technologies, Inc., P. O. Box 6009, Gaithersburg, MD 20897. All Sport vectors contain an
5 ampicillin resistance gene and may be transformed into E. coli strain DH10B, also available from Life Technologies. (See, for instance, Gruber, C. E., et al., Focus 15:59 (1993).) Vector lacmid BA (Bento Soares, Columbia University, NY) contains an ampicillin resistance gene and can be transformed into E. coli strain XL-1 Blue. Vector pCR[®]2.1, which is
10 available from Invitrogen, 1600 Faraday Avenue, Carlsbad, CA 92008, contains an ampicillin resistance gene and may be transformed into E. coli strain DH10B, available from Life Technologies. (See, for instance, Clark, J. M., Nuc. Acids Res. 16:9677-9686 (1988) and Mead, D. et al., Bio/Technology 9: (1991).) Preferably, a polynucleotide of the present invention does not comprise the phage vector sequences identified for the particular clone in Table 1, as well as the corresponding plasmid vector sequences designated above.

15 The deposited material in the sample assigned the ATCC Deposit Number cited in Table 1 for any given cDNA clone also may contain one or more additional plasmids, each comprising a cDNA clone different from that given clone. Thus, deposits sharing the same ATCC Deposit Number contain at least a plasmid for each cDNA clone identified in Table 1. Typically, each ATCC deposit sample cited in Table 1 comprises a mixture of approximately
20 equal amounts (by weight) of about 50 plasmid DNAs, each containing a different cDNA clone; but such a deposit sample may include plasmids for more or less than 50 cDNA clones, up to about 500 cDNA clones.

Two approaches can be used to isolate a particular clone from the deposited sample of plasmid DNAs cited for that clone in Table 1. First, a plasmid is directly isolated by
25 screening the clones using a polynucleotide probe corresponding to SEQ ID NO:X.

Particularly, a specific polynucleotide with 30-40 nucleotides is synthesized using an Applied Biosystems DNA synthesizer according to the sequence reported. The oligonucleotide is labeled, for instance, with ³²P-γ-ATP using T4 polynucleotide kinase and purified according to routine methods. (E.g., Maniatis et al., Molecular Cloning: A
30 Laboratory Manual, Cold Spring Harbor Press, Cold Spring, NY (1982).) The plasmid mixture is transformed into a suitable host, as indicated above (such as XL-1 Blue (Stratagene)) using techniques known to those of skill in the art, such as those provided by

the vector supplier or in related publications or patents cited above. The transformants are plated on 1.5% agar plates (containing the appropriate selection agent, e.g., ampicillin) to a density of about 150 transformants (colonies) per plate. These plates are screened using Nylon membranes according to routine methods for bacterial colony screening (e.g.,
5 Sambrook et al., *Molecular Cloning: A Laboratory Manual*, 2nd Edit., (1989), Cold Spring Harbor Laboratory Press, pages 1.93 to 1.104), or other techniques known to those of skill in the art.

Alternatively, two primers of 17-20 nucleotides derived from both ends of the SEQ ID NO:X (i.e., within the region of SEQ ID NO:X bounded by the 5' NT and the 3' NT of the
10 clone defined in Table 1) are synthesized and used to amplify the desired cDNA using the deposited cDNA plasmid as a template. The polymerase chain reaction is carried out under routine conditions, for instance, in 25 μ l of reaction mixture with 0.5 μ g of the above cDNA template. A convenient reaction mixture is 1.5-5 mM $MgCl_2$, 0.01% (w/v) gelatin, 20 μ M each of dATP, dCTP, dGTP, dTTP, 25 pmol of each primer and 0.25 Unit of Taq
15 polymerase. Thirty five cycles of PCR (denaturation at 94°C for 1 min; annealing at 55°C for 1 min; elongation at 72°C for 1 min) are performed with a Perkin-Elmer Cetus automated thermal cycler. The amplified product is analyzed by agarose gel electrophoresis and the DNA band with expected molecular weight is excised and purified. The PCR product is verified to be the selected sequence by subcloning and sequencing the DNA product.

Several methods are available for the identification of the 5' or 3' non-coding portions of a gene which may not be present in the deposited clone. These methods include but are not limited to, filter probing, clone enrichment using specific probes, and protocols similar or identical to 5' and 3' "RACE" protocols which are well known in the art. For instance, a method similar to 5' RACE is available for generating the missing 5' end of a desired full-
25 length transcript. (Fromont-Racine et al., *Nucleic Acids Res.* 21(7):1683-1684 (1993).)

Briefly, a specific RNA oligonucleotide is ligated to the 5' ends of a population of RNA presumably containing full-length gene RNA transcripts. A primer set containing a primer specific to the ligated RNA oligonucleotide and a primer specific to a known sequence of the gene of interest is used to PCR amplify the 5' portion of the desired full-length gene.

30 This amplified product may then be sequenced and used to generate the full length gene.

This above method starts with total RNA isolated from the desired source, although poly-A⁺ RNA can be used. The RNA preparation can then be treated with phosphatase if

necessary to eliminate 5' phosphate groups on degraded or damaged RNA which may interfere with the later RNA ligase step. The phosphatase should then be inactivated and the RNA treated with tobacco acid pyrophosphatase in order to remove the cap structure present at the 5' ends of messenger RNAs. This reaction leaves a 5' phosphate group at the 5' end of the cap cleaved RNA which can then be ligated to an RNA oligonucleotide using T4 RNA ligase.

This modified RNA preparation is used as a template for first strand cDNA synthesis using a gene specific oligonucleotide. The first strand synthesis reaction is used as a template for PCR amplification of the desired 5' end using a primer specific to the ligated RNA oligonucleotide and a primer specific to the known sequence of the gene of interest. The resultant product is then sequenced and analyzed to confirm that the 5' end sequence belongs to the desired gene.

Example 2: Isolation of Genomic Clones Corresponding to a Polynucleotide

A human genomic P1 library (Genomic Systems, Inc.) is screened by PCR using primers selected for the cDNA sequence corresponding to SEQ ID NO:X., according to the method described in Example 1. (See also, Sambrook.)

Example 3: Tissue specific expression analysis

The Human Genome Sciences, Inc. (HGS) database is derived from sequencing tissue specific cDNA libraries. Libraries generated from a particular tissue are selected and the specific tissue expression pattern of EST groups or assembled contigs within these libraries is determined by comparison of the expression patterns of those groups or contigs within the entire database. ESTs and assembled contigs which show tissue specific expression are selected.

The original clone from which the specific EST sequence was generated, or in the case of an assembled contig, the clone from which the 5' most EST sequence was generated, is obtained from the catalogued library of clones and the insert amplified by PCR using methods known in the art. The PCR product is denatured then transferred in 96 or 384 well format to a nylon membrane (Schleicher and Schuell) generating an array filter of tissue specific clones. Housekeeping genes, maize genes, and known tissue specific genes are included on the filters. These targets can be used in signal normalization and to validate assay

sensitivity. Additional targets are included to monitor probe length and specificity of hybridization.

Radioactively labeled hybridization probes are generated by first strand cDNA synthesis per the manufacturer's instructions (Life Technologies) from mRNA/RNA samples prepared from the specific tissue being analyzed (e.g., prostate, prostate cancer, etc.). The hybridization probes are purified by gel exclusion chromatography, quantitated, and hybridized with the array filters in hybridization bottles at 65°C overnight. The filters are washed under stringent conditions and signals are captured using a Fuji phosphorimager.

Data is extracted using AIS software and following background subtraction, signal normalization is performed. This includes a normalization of filter-wide expression levels between different experimental runs. Genes that are differentially expressed in the tissue of interest are identified.

Example 4: Chromosomal Mapping of the Polynucleotides

An oligonucleotide primer set is designed according to the sequence at the 5' end of SEQ ID NO:X. This primer preferably spans about 100 nucleotides. This primer set is then used in a polymerase chain reaction under the following set of conditions : 30 seconds, 95°C; 1 minute, 56°C; 1 minute, 70°C. This cycle is repeated 32 times followed by one 5 minute cycle at 70°C. Human, mouse, and hamster DNA is used as template in addition to a somatic cell hybrid panel containing individual chromosomes or chromosome fragments (Bios, Inc). The reactions is analyzed on either 8% polyacrylamide gels or 3.5 % agarose gels. Chromosome mapping is determined by the presence of an approximately 100 bp PCR fragment in the particular somatic cell hybrid.

Example 5: Bacterial Expression of a Polypeptide

A polynucleotide encoding a polypeptide of the present invention is amplified using PCR oligonucleotide primers corresponding to the 5' and 3' ends of the DNA sequence, as outlined in Example 1, to synthesize insertion fragments. The primers used to amplify the cDNA insert should preferably contain restriction sites, such as BamHI and XbaI, at the 5' end of the primers in order to clone the amplified product into the expression vector. For example, BamHI and XbaI correspond to the restriction enzyme sites on the bacterial

expression vector pQE-9. (Qiagen, Inc., Chatsworth, CA). This plasmid vector encodes antibiotic resistance (Amp^r), a bacterial origin of replication (ori), an IPTG-regulatable promoter/operator (P/O), a ribosome binding site (RBS), a 6-histidine tag (6-His), and restriction enzyme cloning sites.

5 The pQE-9 vector is digested with BamHI and XbaI and the amplified fragment is ligated into the pQE-9 vector maintaining the reading frame initiated at the bacterial RBS. The ligation mixture is then used to transform the *E. coli* strain M15/rep4 (Qiagen, Inc.) which contains multiple copies of the plasmid pREP4, which expresses the lacI repressor and also confers kanamycin resistance (Kan^r). Transformants are identified by their ability to
10 grow on LB plates and ampicillin/kanamycin resistant colonies are selected. Plasmid DNA is isolated and confirmed by restriction analysis.

Clones containing the desired constructs are grown overnight (O/N) in liquid culture in LB media supplemented with both Amp (100 ug/ml) and Kan (25 ug/ml). The O/N culture is used to inoculate a large culture at a ratio of 1:100 to 1:250. The cells are grown to an
15 optical density 600 (O.D.⁶⁰⁰) of between 0.4 and 0.6. IPTG (Isopropyl-B-D-thiogalactopyranoside) is then added to a final concentration of 1 mM. IPTG induces by inactivating the lacI repressor, clearing the P/O leading to increased gene expression.

Cells are grown for an extra 3 to 4 hours. Cells are then harvested by centrifugation (20 mins at 6000Xg). The cell pellet is solubilized in the chaotropic agent 6 Molar Guanidine
20 HCl by stirring for 3-4 hours at 4°C. The cell debris is removed by centrifugation, and the supernatant containing the polypeptide is loaded onto a nickel-nitrilo-tri-acetic acid ("Ni-NTA") affinity resin column (available from QIAGEN, Inc., *supra*). Proteins with a 6 x His tag bind to the Ni-NTA resin with high affinity and can be purified in a simple one-step procedure (for details see: The QIAexpressionist (1995) QIAGEN, Inc., *supra*).

25 Briefly, the supernatant is loaded onto the column in 6 M guanidine-HCl, pH 8, the column is first washed with 10 volumes of 6 M guanidine-HCl, pH 8, then washed with 10 volumes of 6 M guanidine-HCl pH 6, and finally the polypeptide is eluted with 6 M guanidine-HCl, pH 5.

The purified protein is then renatured by dialyzing it against phosphate-buffered
30 saline (PBS) or 50 mM Na-acetate, pH 6 buffer plus 200 mM NaCl. Alternatively, the protein can be successfully refolded while immobilized on the Ni-NTA column. The

recommended conditions are as follows: renature using a linear 6M-1M urea gradient in 500 mM NaCl, 20% glycerol, 20 mM Tris/HCl pH 7.4, containing protease inhibitors. The renaturation should be performed over a period of 1.5 hours or more. After renaturation the proteins are eluted by the addition of 250 mM imidazole. Imidazole is removed by a final dialyzing step against PBS or 50 mM sodium acetate pH 6 buffer plus 200 mM NaCl. The purified protein is stored at 4° C or frozen at -80° C.

In addition to the above expression vector, the present invention further includes an expression vector comprising phage operator and promoter elements operatively linked to a polynucleotide of the present invention, called pHE4a. (ATCC Accession Number 209645, deposited on February 25, 1998.) This vector contains: 1) a neomycinphosphotransferase gene as a selection marker, 2) an E. coli origin of replication, 3) a T5 phage promoter sequence, 4) two lac operator sequences, 5) a Shine-Delgarno sequence, and 6) the lactose operon repressor gene (lacIq). The origin of replication (oriC) is derived from pUC19 (LTI, Gaithersburg, MD). The promoter sequence and operator sequences are made synthetically.

DNA can be inserted into the pHEa by restricting the vector with NdeI and XbaI, BamHI, XhoI, or Asp718, running the restricted product on a gel, and isolating the larger fragment (the stuffer fragment should be about 310 base pairs). The DNA insert is generated according to the PCR protocol described in Example 1, using PCR primers having restriction sites for NdeI (5' primer) and XbaI, BamHI, XhoI, or Asp718 (3' primer). The PCR insert is gel purified and restricted with compatible enzymes. The insert and vector are ligated according to standard protocols.

The engineered vector could easily be substituted in the above protocol to express protein in a bacterial system.

Example 6: Purification of a Polypeptide from an Inclusion Body

The following alternative method can be used to purify a polypeptide expressed in *E. coli* when it is present in the form of inclusion bodies. Unless otherwise specified, all of the following steps are conducted at 4-10°C.

Upon completion of the production phase of the *E. coli* fermentation, the cell culture is cooled to 4-10°C and the cells harvested by continuous centrifugation at 15,000 rpm (Heraeus Sepatech). On the basis of the expected yield of protein per unit weight of cell paste and the amount of purified protein required, an appropriate amount of cell paste, by

weight, is suspended in a buffer solution containing 100 mM Tris, 50 mM EDTA, pH 7.4. The cells are dispersed to a homogeneous suspension using a high shear mixer.

The cells are then lysed by passing the solution through a microfluidizer (Microfluidics, Corp. or APV Gaulin, Inc.) twice at 4000-6000 psi. The homogenate is then
5 mixed with NaCl solution to a final concentration of 0.5 M NaCl, followed by centrifugation at 7000 xg for 15 min. The resultant pellet is washed again using 0.5M NaCl, 100 mM Tris, 50 mM EDTA, pH 7.4.

The resulting washed inclusion bodies are solubilized with 1.5 M guanidine hydrochloride (GuHCl) for 2-4 hours. After 7000 xg centrifugation for 15 min., the pellet is
10 discarded and the polypeptide containing supernatant is incubated at 4°C overnight to allow further GuHCl extraction.

Following high speed centrifugation (30,000 xg) to remove insoluble particles, the GuHCl solubilized protein is refolded by quickly mixing the GuHCl extract with 20 volumes of buffer containing 50 mM sodium, pH 4.5, 150 mM NaCl, 2 mM EDTA by vigorous
15 stirring. The refolded diluted protein solution is kept at 4°C without mixing for 12 hours prior to further purification steps.

To clarify the refolded polypeptide solution, a previously prepared tangential filtration unit equipped with 0.16 µm membrane filter with appropriate surface area (e.g., Filtron), equilibrated with 40 mM sodium acetate, pH 6.0 is employed. The filtered sample is loaded
20 onto a cation exchange resin (e.g., Poros HS-50, Perseptive Biosystems). The column is washed with 40 mM sodium acetate, pH 6.0 and eluted with 250 mM, 500 mM, 1000 mM, and 1500 mM NaCl in the same buffer, in a stepwise manner. The absorbance at 280 nm of the effluent is continuously monitored. Fractions are collected and further analyzed by SDS-PAGE.

25 Fractions containing the polypeptide are then pooled and mixed with 4 volumes of water. The diluted sample is then loaded onto a previously prepared set of tandem columns of strong anion (Poros HQ-50, Perseptive Biosystems) and weak anion (Poros CM-20, Perseptive Biosystems) exchange resins. The columns are equilibrated with 40 mM sodium acetate, pH 6.0. Both columns are washed with 40 mM sodium acetate, pH 6.0, 200 mM
30 NaCl. The CM-20 column is then eluted using a 10 column volume linear gradient ranging from 0.2 M NaCl, 50 mM sodium acetate, pH 6.0 to 1.0 M NaCl, 50 mM sodium acetate, pH

6.5. Fractions are collected under constant A_{280} monitoring of the effluent. Fractions containing the polypeptide (determined, for instance, by 16% SDS-PAGE) are then pooled.

The resultant polypeptide should exhibit greater than 95% purity after the above refolding and purification steps. No major contaminant bands should be observed from
5 Commassie blue stained 16% SDS-PAGE gel when 5 μ g of purified protein is loaded. The purified protein can also be tested for endotoxin/LPS contamination, and typically the LPS content is less than 0.1 ng/ml according to LAL assays.

Example 7: Cloning and Expression of a Polypeptide in a Baculovirus Expression System

10 In this example, the plasmid shuttle vector pA2 is used to insert a polynucleotide into a baculovirus to express a polypeptide. This expression vector contains the strong polyhedrin promoter of the *Autographa californica* nuclear polyhedrosis virus (AcMNPV) followed by convenient restriction sites such as BamHI, Xba I and Asp718. The polyadenylation site of the simian virus 40 ("SV40") is used for efficient polyadenylation. For easy selection of
15 recombinant virus, the plasmid contains the beta-galactosidase gene from *E. coli* under control of a weak *Drosophila* promoter in the same orientation, followed by the polyadenylation signal of the polyhedrin gene. The inserted genes are flanked on both sides by viral sequences for cell-mediated homologous recombination with wild-type viral DNA to generate a viable virus that express the cloned polynucleotide.

20 Many other baculovirus vectors can be used in place of the vector above, such as pAc373, pVL941, and pAcIM1, as one skilled in the art would readily appreciate, as long as the construct provides appropriately located signals for transcription, translation, secretion and the like, including a signal peptide and an in-frame AUG as required. Such vectors are described, for instance, in Luckow et al., *Virology* 170:31-39 (1989).

25 Specifically, the cDNA sequence contained in ATCC Deposit No.:Z, including the AUG initiation codon, is amplified using the PCR protocol described in Example 1. If a naturally occurring signal sequence is used to produce the polypeptide of the present invention, the pA2 vector does not need a second signal peptide. Alternatively, the vector can be modified (pA2 GP) to include a baculovirus leader sequence, using the standard
30 methods described in Summers et al., "A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures," Texas Agricultural Experimental Station Bulletin No. 1555 (1987).

The amplified fragment is isolated from a 1% agarose gel using a commercially available kit ("GeneClean," BIO 101 Inc., La Jolla, Ca.). The fragment then is digested with appropriate restriction enzymes and again purified on a 1% agarose gel.

The plasmid is digested with the corresponding restriction enzymes and optionally, can be dephosphorylated using calf intestinal phosphatase, using routine procedures known in the art. The DNA is then isolated from a 1% agarose gel using a commercially available kit ("GeneClean" BIO 101 Inc., La Jolla, Ca.).

The fragment and the dephosphorylated plasmid are ligated together with T4 DNA ligase. *E. coli* HB101 or other suitable *E. coli* hosts such as XL-1 Blue (Stratagene Cloning Systems, La Jolla, CA) cells are transformed with the ligation mixture and spread on culture plates. Bacteria containing the plasmid are identified by digesting DNA from individual colonies and analyzing the digestion product by gel electrophoresis. The sequence of the cloned fragment is confirmed by DNA sequencing.

Five µg of a plasmid containing the polynucleotide is co-transfected with 1.0 µg of a commercially available linearized baculovirus DNA ("BaculoGold™ baculovirus DNA", Pharmingen, San Diego, CA), using the lipofection method described by Felgner et al., Proc. Natl. Acad. Sci. USA 84:7413-7417 (1987). One µg of BaculoGold™ virus DNA and 5 µg of the plasmid are mixed in a sterile well of a microtiter plate containing 50 µl of serum-free Grace's medium (Life Technologies Inc., Gaithersburg, MD). Afterwards, 10 µl Lipofectin plus 90 µl Grace's medium are added, mixed and incubated for 15 minutes at room temperature. Then the transfection mixture is added drop-wise to Sf9 insect cells (ATCC CRL 1711) seeded in a 35 mm tissue culture plate with 1 ml Grace's medium without serum. The plate is then incubated for 5 hours at 27° C. The transfection solution is then removed from the plate and 1 ml of Grace's insect medium supplemented with 10% fetal calf serum is added. Cultivation is then continued at 27° C for four days.

After four days the supernatant is collected and a plaque assay is performed, as described by Summers and Smith, *supra*. An agarose gel with "Blue Gal" (Life Technologies Inc., Gaithersburg) is used to allow easy identification and isolation of gal-expressing clones, which produce blue-stained plaques. (A detailed description of a "plaque assay" of this type can also be found in the user's guide for insect cell culture and baculovirology distributed by Life Technologies Inc., Gaithersburg, page 9-10.) After appropriate incubation, blue stained plaques are picked with the tip of a micropipettor (e.g., Eppendorf). The agar containing the

recombinant viruses is then resuspended in a microcentrifuge tube containing 200 μ l of Grace's medium and the suspension containing the recombinant baculovirus is used to infect Sf9 cells seeded in 35 mm dishes. Four days later the supernatants of these culture dishes are harvested and then they are stored at 4° C.

5 To verify the expression of the polypeptide, Sf9 cells are grown in Grace's medium supplemented with 10% heat-inactivated FBS. The cells are infected with the recombinant baculovirus containing the polynucleotide at a multiplicity of infection ("MOI") of about 2. If radiolabeled proteins are desired, 6 hours later the medium is removed and is replaced with SF900 II medium minus methionine and cysteine (available from Life Technologies Inc.,
10 Rockville, MD). After 42 hours, 5 μ Ci of 35 S-methionine and 5 μ Ci 35 S-cysteine (available from Amersham) are added. The cells are further incubated for 16 hours and then are harvested by centrifugation. The proteins in the supernatant as well as the intracellular proteins are analyzed by SDS-PAGE followed by autoradiography (if radiolabeled).

Microsequencing of the amino acid sequence of the amino terminus of purified
15 protein may be used to determine the amino terminal sequence of the produced protein.

Example 8: Expression of a Polypeptide in Mammalian Cells

The polypeptide of the present invention can be expressed in a mammalian cell. A typical mammalian expression vector contains a promoter element, which mediates the
20 initiation of transcription of mRNA, a protein coding sequence, and signals required for the termination of transcription and polyadenylation of the transcript. Additional elements include enhancers, Kozak sequences and intervening sequences flanked by donor and acceptor sites for RNA splicing. Highly efficient transcription is achieved with the early and late promoters from SV40, the long terminal repeats (LTRs) from Retroviruses, e.g., RSV,
25 HTLV1, HIV1 and the early promoter of the cytomegalovirus (CMV). However, cellular elements can also be used (e.g., the human actin promoter).

Suitable expression vectors for use in practicing the present invention include, for example, vectors such as pSVL and pMSG (Pharmacia, Uppsala, Sweden), pRSVcat (ATCC 37152), pSV2dhfr (ATCC 37146), pBC12MI (ATCC 67109), pCMVSPORT 2.0, and
30 pCMVSPORT 3.0. Mammalian host cells that could be used include, human Hela, 293, H9 and Jurkat cells, mouse NIH3T3 and C127 cells, Cos 1, Cos 7 and CV1, quail QC1-3 cells, mouse L cells and Chinese hamster ovary (CHO) cells.

Alternatively, the polypeptide can be expressed in stable cell lines containing the polynucleotide integrated into a chromosome. The co-transfection with a selectable marker such as DHFR, gpt, neomycin, hygromycin allows the identification and isolation of the transfected cells.

5 The transfected gene can also be amplified to express large amounts of the encoded protein. The DHFR (dihydrofolate reductase) marker is useful in developing cell lines that carry several hundred or even several thousand copies of the gene of interest. (See, e.g., Alt, F. W., et al., *J. Biol. Chem.* 253:1357-1370 (1978); Hamlin, J. L. and Ma, C., *Biochem. et Biophys. Acta*, 1097:107-143 (1990); Page, M. J. and Sydenham, M. A., *Biotechnology* 9:64-10 68 (1991).) Another useful selection marker is the enzyme glutamine synthase (GS) (Murphy et al., *Biochem J.* 227:277-279 (1991); Bebbington et al., *Bio/Technology* 10:169-175 (1992). Using these markers, the mammalian cells are grown in selective medium and the cells with the highest resistance are selected. These cell lines contain the amplified gene(s) integrated into a chromosome. Chinese hamster ovary (CHO) and NSO cells are often used
15 for the production of proteins.

Derivatives of the plasmid pSV2-dhfr (ATCC Accession No. 37146), the expression vectors pC4 (ATCC Accession No. 209646) and pC6 (ATCC Accession No. 209647) contain the strong promoter (LTR) of the Rous Sarcoma Virus (Cullen et al., *Molecular and Cellular Biology*, 438-447 (March, 1985)) plus a fragment of the CMV-enhancer (Boshart et al., *Cell* 20 41:521-530 (1985).) Multiple cloning sites, e.g., with the restriction enzyme cleavage sites BamHI, XbaI and Asp718, facilitate the cloning of the gene of interest. The vectors also contain the 3' intron, the polyadenylation and termination signal of the rat preproinsulin gene, and the mouse DHFR gene under control of the SV40 early promoter.

Specifically, the plasmid pC6, for example, is digested with appropriate restriction
25 enzymes and then dephosphorylated using calf intestinal phosphates by procedures known in the art. The vector is then isolated from a 1% agarose gel.

A polynucleotide of the present invention is amplified according to the protocol outlined in Example 1. If a naturally occurring signal sequence is used to produce the polypeptide of the present invention, the vector does not need a second signal peptide.
30 Alternatively, if a naturally occurring signal sequence is not used, the vector can be modified to include a heterologous signal sequence. (See, e.g., WO 96/34891.)

The amplified fragment is isolated from a 1% agarose gel using a commercially available kit ("GeneClean," BIO 101 Inc., La Jolla, Ca.). The fragment then is digested with appropriate restriction enzymes and again purified on a 1% agarose gel.

The amplified fragment is then digested with the same restriction enzyme and purified
5 on a 1% agarose gel. The isolated fragment and the dephosphorylated vector are then ligated with T4 DNA ligase. *E. coli* HB101 or XL-1 Blue cells are then transformed and bacteria are identified that contain the fragment inserted into plasmid pC6 using, for instance, restriction enzyme analysis.

Chinese hamster ovary cells lacking an active DHFR gene is used for transfection.

10 Five μ g of the expression plasmid pC6 or pC4 is cotransfected with 0.5 μ g of the plasmid pSVneo using lipofectin (Felgner et al., *supra*). The plasmid pSV2-neo contains a dominant selectable marker, the *neo* gene from Tn5 encoding an enzyme that confers resistance to a group of antibiotics including G418. The cells are seeded in alpha minus MEM supplemented with 1 mg/ml G418. After 2 days, the cells are trypsinized and seeded in
15 hybridoma cloning plates (Greiner, Germany) in alpha minus MEM supplemented with 10, 25, or 50 ng/ml of methotrexate plus 1 mg/ml G418. After about 10-14 days single clones are trypsinized and then seeded in 6-well petri dishes or 10 ml flasks using different concentrations of methotrexate (50 nM, 100 nM, 200 nM, 400 nM, 800 nM). Clones growing at the highest concentrations of methotrexate are then transferred to new 6-well plates
20 containing even higher concentrations of methotrexate (1 μ M, 2 μ M, 5 μ M, 10 mM, 20 mM). The same procedure is repeated until clones are obtained which grow at a concentration of 100 - 200 μ M. Expression of the desired gene product is analyzed, for instance, by SDS-PAGE and Western blot or by reversed phase HPLC analysis.

25 **Example 9: Protein Fusions**

The polypeptides of the present invention are preferably fused to other proteins. These fusion proteins can be used for a variety of applications. For example, fusion of the present polypeptides to His-tag, HA-tag, protein A, IgG domains, and maltose binding protein facilitates purification. (See Example 5; see also EP A 394,827; Traunecker, et al.,
30 Nature 331:84-86 (1988).) Similarly, fusion to IgG-1, IgG-3, and albumin increases the halflife time in vivo. Nuclear localization signals fused to the polypeptides of the present invention can target the protein to a specific subcellular localization, while covalent

heterodimer or homodimers can increase or decrease the activity of a fusion protein. Fusion proteins can also create chimeric molecules having more than one function. Finally, fusion proteins can increase solubility and/or stability of the fused protein compared to the non-fused protein. All of the types of fusion proteins described above can be made by modifying the following protocol, which outlines the fusion of a polypeptide to an IgG molecule, or the protocol described in Example 5.

Briefly, the human Fc portion of the IgG molecule can be PCR amplified, using primers that span the 5' and 3' ends of the sequence described below. These primers also should have convenient restriction enzyme sites that will facilitate cloning into an expression vector, preferably a mammalian expression vector.

For example, if pC4 (Accession No. 209646) is used, the human Fc portion can be ligated into the BamHI cloning site. Note that the 3' BamHI site should be destroyed. Next, the vector containing the human Fc portion is re-restricted with BamHI, linearizing the vector, and a polynucleotide of the present invention, isolated by the PCR protocol described in Example 1, is ligated into this BamHI site. Note that the polynucleotide is cloned without a stop codon, otherwise a fusion protein will not be produced.

If the naturally occurring signal sequence is used to produce the polypeptide of the present invention, pC4 does not need a second signal peptide. Alternatively, if the naturally occurring signal sequence is not used, the vector can be modified to include a heterologous signal sequence. (See, e.g., WO 96/34891.)

Human IgG Fc region:

```
GGGATCCGGAGCCCAAATCTTCTGACAAACTCACACATGCCCACCGTGCCCAG
CACCTGAATTTCGAGGGTGCACCGTCAGTCTTCTCTTCCCCCAAACCCAAGGA
CACCTCATGATCTCCCGGACTCCTGAGGTCACATGCGTGGTGGTGGACGTAAGC
CACGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCAT
AATGCCAAGACAAAGCCGCGGGAGGAGCAGTACAACAGCACGTACCGTGTGGTC
AGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGC
AAGGTCTCCAACAAAGCCCTCCCAACCCCCATCGAGAAAACCATCTCCAAAGCC
AAAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGATGAG
CTGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCAAGCG
ACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACACTACAAGACC
```

ACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCG
TGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTCCGTGATGCATG
AGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATG
AGTGCGACGGCCGCGACTCTAGAGGAT (SEQ ID NO:1)

5

Example 10: Production of an Antibody from a Polypeptide

a) Hybridoma Technology

The antibodies of the present invention can be prepared by a variety of methods. (See, Current Protocols, Chapter 2.) As one example of such methods, cells expressing polypeptide of the present invention are administered to an animal to induce the production of sera containing polyclonal antibodies. In a preferred method, a preparation of polypeptide of the present invention is prepared and purified to render it substantially free of natural contaminants. Such a preparation is then introduced into an animal in order to produce polyclonal antisera of greater specific activity.

Monoclonal antibodies specific for polypeptide of the present invention are prepared using hybridoma technology. (Kohler et al., Nature 256:495 (1975); Kohler et al., Eur. J. Immunol. 6:511 (1976); Kohler et al., Eur. J. Immunol. 6:292 (1976); Hammerling et al., in: Monoclonal Antibodies and T-Cell Hybridomas, Elsevier, N.Y., pp. 563-681 (1981)). In general, an animal (preferably a mouse) is immunized with polypeptide of the present invention or, more preferably, with a secreted polypeptide of the present invention-expressing cell. Such polypeptide-expressing cells are cultured in any suitable tissue culture medium, preferably in Earle's modified Eagle's medium supplemented with 10% fetal bovine serum (inactivated at about 56°C), and supplemented with about 10 g/l of nonessential amino acids, about 1,000 U/ml of penicillin, and about 100 µg/ml of streptomycin.

The splenocytes of such mice are extracted and fused with a suitable myeloma cell line. Any suitable myeloma cell line may be employed in accordance with the present invention; however, it is preferable to employ the parent myeloma cell line (SP2O), available from the ATCC. After fusion, the resulting hybridoma cells are selectively maintained in HAT medium, and then cloned by limiting dilution as described by Wands et al. (Gastroenterology 80:225-232 (1981)). The hybridoma cells obtained through such a selection are then assayed to identify clones which secrete antibodies capable of binding the polypeptide of the present invention.

Alternatively, additional antibodies capable of binding to polypeptide of the present invention can be produced in a two-step procedure using anti-idiotypic antibodies. Such a method makes use of the fact that antibodies are themselves antigens, and therefore, it is possible to obtain an antibody which binds to a second antibody. In accordance with this method, protein specific antibodies are used to immunize an animal, preferably a mouse. The splenocytes of such an animal are then used to produce hybridoma cells, and the hybridoma cells are screened to identify clones which produce an antibody whose ability to bind to the polypeptide of the present invention-specific antibody can be blocked by polypeptide of the present invention. Such antibodies comprise anti-idiotypic antibodies to the polypeptide of the present invention-specific antibody and are used to immunize an animal to induce formation of further polypeptide of the present invention-specific antibodies.

For in vivo use of antibodies in humans, an antibody is "humanized". Such antibodies can be produced using genetic constructs derived from hybridoma cells producing the monoclonal antibodies described above. Methods for producing chimeric and humanized antibodies are known in the art and are discussed herein. (See, for review, Morrison, Science 229:1202 (1985); Oi et al., BioTechniques 4:214 (1986); Cabilly et al., U.S. Patent No. 4,816,567; Taniguchi et al., EP 171496; Morrison et al., EP 173494; Neuberger et al., WO 8601533; Robinson et al., WO 8702671; Boulianne et al., Nature 312:643 (1984); Neuberger et al., Nature 314:268 (1985).)

b) Isolation Of Antibody Fragments Directed Against Polypeptide of the Present Invention From A Library Of scFvs

Naturally occurring V-genes isolated from human PBLs are constructed into a library of antibody fragments which contain reactivities against polypeptide of the present invention to which the donor may or may not have been exposed (see e.g., U.S. Patent 5,885,793 incorporated herein by reference in its entirety).

Rescue of the Library. A library of scFvs is constructed from the RNA of human PBLs as described in PCT publication WO 92/01047. To rescue phage displaying antibody fragments, approximately 10⁹ E. coli harboring the phagemid are used to inoculate 50 ml of 2xTY containing 1% glucose and 100 µg/ml of ampicillin (2xTY-AMP-GLU) and grown to an O.D. of 0.8 with shaking. Five ml of this culture is used to innoculate 50 ml of 2xTY-AMP-GLU, 2 x 10⁸ TU of delta gene 3 helper (M13 delta gene III, see PCT publication WO

92/01047) are added and the culture incubated at 37°C for 45 minutes without shaking and then at 37°C for 45 minutes with shaking. The culture is centrifuged at 4000 r.p.m. for 10 min. and the pellet resuspended in 2 liters of 2xTY containing 100 µg/ml ampicillin and 50 µg/ml kanamycin and grown overnight. Phage are prepared as described in PCT publication
5 WO 92/01047.

M13 delta gene III is prepared as follows: M13 delta gene III helper phage does not encode gene III protein, hence the phage(mid) displaying antibody fragments have a greater avidity of binding to antigen. Infectious M13 delta gene III particles are made by growing the helper phage in cells harboring a pUC19 derivative supplying the wild type gene III
10 protein during phage morphogenesis. The culture is incubated for 1 hour at 37° C without shaking and then for a further hour at 37°C with shaking. Cells are spun down (IEC-Centra 8,400 r.p.m. for 10 min), resuspended in 300 ml 2xTY broth containing 100 µg ampicillin/ml and 25 µg kanamycin/ml (2xTY-AMP-KAN) and grown overnight, shaking at 37°C. Phage particles are purified and concentrated from the culture medium by two PEG-precipitations
15 (Sambrook et al., 1990), resuspended in 2 ml PBS and passed through a 0.45 µm filter (Minisart NML; Sartorius) to give a final concentration of approximately 10¹³ transducing units/ml (ampicillin-resistant clones).

Panning of the Library. Immunotubes (Nunc) are coated overnight in PBS with 4 ml of either 100 µg/ml or 10 µg/ml of a polypeptide of the present invention. Tubes are blocked
20 with 2% Marvel-PBS for 2 hours at 37°C and then washed 3 times in PBS. Approximately 10¹³ TU of phage is applied to the tube and incubated for 30 minutes at room temperature tumbling on an over and under turntable and then left to stand for another 1.5 hours. Tubes are washed 10 times with PBS 0.1% Tween-20 and 10 times with PBS. Phage are eluted by adding 1 ml of 100 mM triethylamine and rotating 15 minutes on an under and over turntable
25 after which the solution is immediately neutralized with 0.5 ml of 1.0M Tris-HCl, pH 7.4. Phage are then used to infect 10 ml of mid-log E. coli TG1 by incubating eluted phage with bacteria for 30 minutes at 37°C. The E. coli are then plated on TYE plates containing 1% glucose and 100 µg/ml ampicillin. The resulting bacterial library is then rescued with delta gene 3 helper phage as described above to prepare phage for a subsequent round of selection.
30 This process is then repeated for a total of 4 rounds of affinity purification with tube-washing increased to 20 times with PBS, 0.1% Tween-20 and 20 times with PBS for rounds 3 and 4.

Characterization of Binders. Eluted phage from the 3rd and 4th rounds of selection

are used to infect *E. coli* HB 2151 and soluble scFv is produced (Marks, et al., 1991) from single colonies for assay. ELISAs are performed with microtitre plates coated with either 10 pg/ml of the polypeptide of the present invention in 50 mM bicarbonate pH 9.6. Clones positive in ELISA are further characterized by PCR fingerprinting (see, e.g., PCT publication
5 WO 92/01047) and then by sequencing. These ELISA positive clones may also be further characterized by techniques known in the art, such as, for example, epitope mapping, binding affinity, receptor signal transduction, ability to block or competitively inhibit antibody/antigen binding, and competitive agonistic or antagonistic activity.

10 ***Example 11: Method of Determining Alterations in a Gene Corresponding to a Polynucleotide***

RNA isolated from entire families or individual patients presenting with a phenotype of interest (such as a disease) is be isolated. cDNA is then generated from these RNA samples using protocols known in the art. (See, Sambrook.) The cDNA is then used as a
15 template for PCR, employing primers surrounding regions of interest in SEQ ID NO:X and/or the nucleotide sequence of a deposited cDNA clone referenced in Table 1. Suggested PCR conditions consist of 35 cycles at 95 degrees C for 30 seconds; 60-120 seconds at 52-58 degrees C; and 60-120 seconds at 70 degrees C, using buffer solutions described in Sidransky et al., Science 252:706 (1991).

20 PCR products are then sequenced using primers labeled at their 5' end with T4 polynucleotide kinase, employing SequiTherm Polymerase. (Epicentre Technologies). The intron-exon borders of selected exons is also determined and genomic PCR products analyzed to confirm the results. PCR products harboring suspected mutations is then cloned and sequenced to validate the results of the direct sequencing.

25 PCR products is cloned into T-tailed vectors as described in Holton et al., Nucleic Acids Research, 19:1156 (1991) and sequenced with T7 polymerase (United States Biochemical). Affected individuals are identified by mutations not present in unaffected individuals.

30 Genomic rearrangements are also observed as a method of determining alterations in a gene corresponding to a polynucleotide. Genomic clones isolated according to Example 2 are nick-translated with digoxigenindeoxy-uridine 5'-triphosphate (Boehringer Mannheim), and FISH performed as described in Johnson et al., Methods Cell Biol. 35:73-99 (1991).

Hybridization with the labeled probe is carried out using a vast excess of human cot-1 DNA for specific hybridization to the corresponding genomic locus.

Chromosomes are counterstained with 4,6-diamino-2-phenylidole and propidium iodide, producing a combination of C- and R-bands. Aligned images for precise mapping are obtained using a triple-band filter set (Chroma Technology, Brattleboro, VT) in combination with a cooled charge-coupled device camera (Photometrics, Tucson, AZ) and variable excitation wavelength filters. (Johnson et al., Genet. Anal. Tech. Appl., 8:75 (1991).) Image collection, analysis and chromosomal fractional length measurements are performed using the ISee Graphical Program System. (Inovision Corporation, Durham, NC.) Chromosome alterations of the genomic region hybridized by the probe are identified as insertions, deletions, and translocations. These alterations are used as a diagnostic marker for an associated disease.

Example 12: Method of Detecting Abnormal Levels of a Polypeptide in a Biological Sample

A polypeptide of the present invention can be detected in a biological sample, and if an increased or decreased level of the polypeptide is detected, this polypeptide is a marker for a particular phenotype. Methods of detection are numerous, and thus, it is understood that one skilled in the art can modify the following assay to fit their particular needs.

For example, antibody-sandwich ELISAs are used to detect polypeptides in a sample, preferably a biological sample. Wells of a microtiter plate are coated with specific antibodies, at a final concentration of 0.2 to 10 ug/ml. The antibodies are either monoclonal or polyclonal and are produced by the method described in Example 10. The wells are blocked so that non-specific binding of the polypeptide to the well is reduced.

The coated wells are then incubated for > 2 hours at RT with a sample containing the polypeptide. Preferably, serial dilutions of the sample should be used to validate results. The plates are then washed three times with deionized or distilled water to remove unbounded polypeptide.

Next, 50 ul of specific antibody-alkaline phosphatase conjugate, at a concentration of 25-400 ng, is added and incubated for 2 hours at room temperature. The plates are again washed three times with deionized or distilled water to remove unbounded conjugate.

Add 75 ul of 4-methylumbelliferyl phosphate (MUP) or p-nitrophenyl phosphate (NPP) substrate solution to each well and incubate 1 hour at room temperature. Measure the reaction by a microtiter plate reader. Prepare a standard curve, using serial dilutions of a control sample, and plot polypeptide concentration on the X-axis (log scale) and fluorescence or absorbance of the Y-axis (linear scale). Interpolate the concentration of the polypeptide in the sample using the standard curve.

Example 13: Formulation

The invention also provides methods of treatment and/or prevention of diseases or disorders (such as, for example, any one or more of the diseases or disorders disclosed herein) by administration to a subject of an effective amount of a Therapeutic. By therapeutic is meant a polynucleotides or polypeptides of the invention (including fragments and variants), agonists or antagonists thereof, and/or antibodies thereto, in combination with a pharmaceutically acceptable carrier type (e.g., a sterile carrier).

The Therapeutic will be formulated and dosed in a fashion consistent with good medical practice, taking into account the clinical condition of the individual patient (especially the side effects of treatment with the Therapeutic alone), the site of delivery, the method of administration, the scheduling of administration, and other factors known to practitioners. The "effective amount" for purposes herein is thus determined by such considerations.

As a general proposition, the total pharmaceutically effective amount of the Therapeutic administered parenterally per dose will be in the range of about 1ug/kg/day to 10 mg/kg/day of patient body weight, although, as noted above, this will be subject to therapeutic discretion. More preferably, this dose is at least 0.01 mg/kg/day, and most preferably for humans between about 0.01 and 1 mg/kg/day for the hormone. If given continuously, the Therapeutic is typically administered at a dose rate of about 1 ug/kg/hour to about 50 ug/kg/hour, either by 1-4 injections per day or by continuous subcutaneous infusions, for example, using a mini-pump. An intravenous bag solution may also be employed. The length of treatment needed to observe changes and the interval following treatment for responses to occur appears to vary depending on the desired effect.

Therapeutics can be administered orally, rectally, parenterally, intracisternally, intravaginally, intraperitoneally, topically (as by powders, ointments, gels, drops or

transdermal patch), buccally, or as an oral or nasal spray. "Pharmaceutically acceptable carrier" refers to a non-toxic solid, semisolid or liquid filler, diluent, encapsulating material or formulation auxiliary of any. The term "parenteral" as used herein refers to modes of administration which include intravenous, intramuscular, intraperitoneal, intrasternal, subcutaneous and intraarticular injection and infusion.

Therapeutics of the invention are also suitably administered by sustained-release systems. Suitable examples of sustained-release Therapeutics are administered orally, rectally, parenterally, intracisternally, intravaginally, intraperitoneally, topically (as by powders, ointments, gels, drops or transdermal patch), buccally, or as an oral or nasal spray.

"Pharmaceutically acceptable carrier" refers to a non-toxic solid, semisolid or liquid filler, diluent, encapsulating material or formulation auxiliary of any type. The term "parenteral" as used herein refers to modes of administration which include intravenous, intramuscular, intraperitoneal, intrasternal, subcutaneous and intraarticular injection and infusion.

Therapeutics of the invention are also suitably administered by sustained-release systems. Suitable examples of sustained-release Therapeutics include suitable polymeric materials (such as, for example, semi-permeable polymer matrices in the form of shaped articles, e.g., films, or microcapsules), suitable hydrophobic materials (for example as an emulsion in an acceptable oil) or ion exchange resins, and sparingly soluble derivatives (such as, for example, a sparingly soluble salt).

Sustained-release matrices include polylactides (U.S. Pat. No. 3,773,919, EP 58,481), copolymers of L-glutamic acid and gamma-ethyl-L-glutamate (Sidman et al., *Biopolymers* 22:547-556 (1983)), poly (2- hydroxyethyl methacrylate) (Langer et al., *J. Biomed. Mater. Res.* 15:167-277 (1981), and Langer, *Chem. Tech.* 12:98-105 (1982)), ethylene vinyl acetate (Langer et al., *Id.*) or poly-D- (-)-3-hydroxybutyric acid (EP 133,988).

Sustained-release Therapeutics also include liposomally entrapped Therapeutics of the invention (*see generally*, Langer, *Science* 249:1527-1533 (1990); Treat et al., in *Liposomes in the Therapy of Infectious Disease and Cancer*, Lopez-Berestein and Fidler (eds.), Liss, New York, pp. 317 -327 and 353-365 (1989)). Liposomes containing the Therapeutic are prepared by methods known per se: DE 3,218,121; Epstein et al., *Proc. Natl. Acad. Sci. (USA)* 82:3688-3692 (1985); Hwang et al., *Proc. Natl. Acad. Sci. (USA)* 77:4030-4034 (1980); EP 52,322; EP 36,676; EP 88,046; EP 143,949; EP 142,641; Japanese Pat. Appl. 83-118008; U.S. Pat. Nos. 4,485,045 and 4,544,545; and EP 102,324. Ordinarily, the

liposomes are of the small (about 200-800 Angstroms) unilamellar type in which the lipid content is greater than about 30 mol. percent cholesterol, the selected proportion being adjusted for the optimal Therapeutic.

In yet an additional embodiment, the Therapeutics of the invention are delivered by way of a pump (*see* Langer, *supra*; Sefton, CRC Crit. Ref. Biomed. Eng. 14:201 (1987); Buchwald et al., Surgery 88:507 (1980); Saudek et al., N. Engl. J. Med. 321:574 (1989)).

Other controlled release systems are discussed in the review by Langer (*Science* 249:1527-1533 (1990)).

For parenteral administration, in one embodiment, the Therapeutic is formulated generally by mixing it at the desired degree of purity, in a unit dosage injectable form (solution, suspension, or emulsion), with a pharmaceutically acceptable carrier, i.e., one that is non-toxic to recipients at the dosages and concentrations employed and is compatible with other ingredients of the formulation. For example, the formulation preferably does not include oxidizing agents and other compounds that are known to be deleterious to the Therapeutic.

Generally, the formulations are prepared by contacting the Therapeutic uniformly and intimately with liquid carriers or finely divided solid carriers or both. Then, if necessary, the product is shaped into the desired formulation. Preferably the carrier is a parenteral carrier, more preferably a solution that is isotonic with the blood of the recipient. Examples of such carrier vehicles include water, saline, Ringer's solution, and dextrose solution. Non-aqueous vehicles such as fixed oils and ethyl oleate are also useful herein, as well as liposomes.

The carrier suitably contains minor amounts of additives such as substances that enhance isotonicity and chemical stability. Such materials are non-toxic to recipients at the dosages and concentrations employed, and include buffers such as phosphate, citrate, succinate, acetic acid, and other organic acids or their salts; antioxidants such as ascorbic acid; low molecular weight (less than about ten residues) polypeptides, e.g., polyarginine or tripeptides; proteins, such as serum albumin, gelatin, or immunoglobulins; hydrophilic polymers such as polyvinylpyrrolidone; amino acids, such as glycine, glutamic acid, aspartic acid, or arginine; monosaccharides, disaccharides, and other carbohydrates including cellulose or its derivatives, glucose, manose, or dextrans; chelating agents such as EDTA; sugar alcohols such as mannitol or sorbitol; counterions such as sodium; and/or nonionic surfactants such as polysorbates, poloxamers, or PEG.

The Therapeutic is typically formulated in such vehicles at a concentration of about 0.1 mg/ml to 100 mg/ml, preferably 1-10 mg/ml, at a pH of about 3 to 8. It will be understood that the use of certain of the foregoing excipients, carriers, or stabilizers will result in the formation of polypeptide salts.

5 Any pharmaceutical used for therapeutic administration can be sterile. Sterility is readily accomplished by filtration through sterile filtration membranes (e.g., 0.2 micron membranes). Therapeutics generally are placed into a container having a sterile access port, for example, an intravenous solution bag or vial having a stopper pierceable by a hypodermic injection needle.

10 Therapeutics ordinarily will be stored in unit or multi-dose containers, for example, sealed ampoules or vials, as an aqueous solution or as a lyophilized formulation for reconstitution. As an example of a lyophilized formulation, 10-ml vials are filled with 5 ml of sterile-filtered 1% (w/v) aqueous Therapeutic solution, and the resulting mixture is lyophilized. The infusion solution is prepared by reconstituting the lyophilized Therapeutic
15 using bacteriostatic Water-for-Injection.

The invention also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the Therapeutics of the invention. Associated with such container(s) can be a notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of pharmaceuticals or biological products,
20 which notice reflects approval by the agency of manufacture, use or sale for human administration. In addition, the Therapeutics may be employed in conjunction with other therapeutic compounds.

The Therapeutics of the invention may be administered alone or in combination with adjuvants. Adjuvants that may be administered with the Therapeutics of the invention
25 include, but are not limited to, alum, alum plus deoxycholate (ImmunoAg), MTP-PE (Biocine Corp.), QS21 (Genentech, Inc.), BCG, and MPL. In a specific embodiment, Therapeutics of the invention are administered in combination with alum. In another specific embodiment, Therapeutics of the invention are administered in combination with QS-21. Further adjuvants that may be administered with the Therapeutics of the invention include, but are not limited
30 to, Monophosphoryl lipid immunomodulator, AdjuVax 100a, QS-21, QS-18, CRL1005, Aluminum salts, MF-59, and Virosomal adjuvant technology. Vaccines that may be administered with the Therapeutics of the invention include, but are not limited to, vaccines

directed toward protection against MMR (measles, mumps, rubella), polio, varicella, tetanus/diphtheria, hepatitis A, hepatitis B, haemophilus influenzae B, whooping cough, pneumonia, influenza, Lyme's Disease, rotavirus, cholera, yellow fever, Japanese encephalitis, poliomyelitis, rabies, typhoid fever, and pertussis. Combinations may be administered either concomitantly, e.g., as an admixture, separately but simultaneously or concurrently; or sequentially. This includes presentations in which the combined agents are administered together as a therapeutic mixture, and also procedures in which the combined agents are administered separately but simultaneously, e.g., as through separate intravenous lines into the same individual. Administration "in combination" further includes the separate administration of one of the compounds or agents given first, followed by the second.

The Therapeutics of the invention may be administered alone or in combination with other therapeutic agents. Therapeutic agents that may be administered in combination with the Therapeutics of the invention, include but not limited to, other members of the TNF family, chemotherapeutic agents, antibiotics, steroidal and non-steroidal anti-inflammatories, conventional immunotherapeutic agents, cytokines and/or growth factors. Combinations may be administered either concomitantly, e.g., as an admixture, separately but simultaneously or concurrently; or sequentially. This includes presentations in which the combined agents are administered together as a therapeutic mixture, and also procedures in which the combined agents are administered separately but simultaneously, e.g., as through separate intravenous lines into the same individual. Administration "in combination" further includes the separate administration of one of the compounds or agents given first, followed by the second.

In one embodiment, the Therapeutics of the invention are administered in combination with members of the TNF family. TNF, TNF-related or TNF-like molecules that may be administered with the Therapeutics of the invention include, but are not limited to, soluble forms of TNF-alpha, lymphotoxin-alpha (LT-alpha, also known as TNF-beta), LT-beta (found in complex heterotrimer LT-alpha2-beta), OPGL, FasL, CD27L, CD30L, CD40L, 4-1BBL, DcR3, OX40L, TNF-gamma (International Publication No. WO 96/14328), AIM-I (International Publication No. WO 97/33899), endokine-alpha (International Publication No. WO 98/07880), TR6 (International Publication No. WO 98/30694), OPG, and neutrokin-alpha (International Publication No. WO 98/18921, OX40, and nerve growth factor (NGF), and soluble forms of Fas, CD30, CD27, CD40 and 4-1BB, TR2 (International Publication No. WO 96/34095), DR3 (International Publication No. WO 97/33904), DR4

(International Publication No. WO 98/32856), TR5 (International Publication No. WO 98/30693), TR6 (International Publication No. WO 98/30694), TR7 (International Publication No. WO 98/41629), TRANK, TR9 (International Publication No. WO 98/56892), TR10 (International Publication No. WO 98/54202), 312C2 (International Publication No. WO 98/06842), and TR12, and soluble forms CD154, CD70, and CD153.

In certain embodiments, Therapeutics of the invention are administered in combination with antiretroviral agents, nucleoside reverse transcriptase inhibitors, non-nucleoside reverse transcriptase inhibitors, and/or protease inhibitors. Nucleoside reverse transcriptase inhibitors that may be administered in combination with the Therapeutics of the invention, include, but are not limited to, RETROVIR™ (zidovudine/AZT), VIDEX™ (didanosine/ddI), HIVID™ (zalcitabine/ddC), ZERIT™ (stavudine/d4T), EPIVIR™ (lamivudine/3TC), and COMBIVIR™ (zidovudine/lamivudine). Non-nucleoside reverse transcriptase inhibitors that may be administered in combination with the Therapeutics of the invention, include, but are not limited to, VIRAMUNE™ (nevirapine), RESCRIPTOR™ (delavirdine), and SUSTIVA™ (efavirenz). Protease inhibitors that may be administered in combination with the Therapeutics of the invention, include, but are not limited to, CRIXIVAN™ (indinavir), NORVIR™ (ritonavir), INVIRASE™ (saquinavir), and VIRACEPT™ (nelfinavir). In a specific embodiment, antiretroviral agents, nucleoside reverse transcriptase inhibitors, non-nucleoside reverse transcriptase inhibitors, and/or protease inhibitors may be used in any combination with Therapeutics of the invention to treat AIDS and/or to prevent or treat HIV infection.

In other embodiments, Therapeutics of the invention may be administered in combination with anti-opportunistic infection agents. Anti-opportunistic agents that may be administered in combination with the Therapeutics of the invention, include, but are not limited to, TRIMETHOPRIM-SULFAMETHOXAZOLE™, DAPSONE™, PENTAMIDINE™, ATOVAQUONE™, ISONIAZID™, RIFAMPIN™, PYRAZINAMIDE™, ETHAMBUTOL™, RIFABUTIN™, CLARITHROMYCIN™, AZITHROMYCIN™, GANCICLOVIR™, FOSCARNET™, CIDOFOVIR™, FLUCONAZOLE™, ITRACONAZOLE™, KETOCONAZOLE™, ACYCLOVIR™, FAMCICLOVIR™, PYRIMETHAMINE™, LEUCOVORIN™, NEUPOGEN™ (filgrastim/G-CSF), and LEUKINE™ (sargramostim/GM-CSF). In a specific embodiment, Therapeutics of the

invention are used in any combination with TRIMETHOPRIM-SULFAMETHOXAZOLE™, DAPSONE™, PENTAMIDINE™, and/or ATOVAQUONE™ to prophylactically treat or prevent an opportunistic *Pneumocystis carinii* pneumonia infection. In another specific embodiment, Therapeutics of the invention are used in any combination with ISONIAZID™, RIFAMPIN™, PYRAZINAMIDE™, and/or ETHAMBUTOL™ to prophylactically treat or prevent an opportunistic *Mycobacterium avium* complex infection. In another specific embodiment, Therapeutics of the invention are used in any combination with RIFABUTIN™, CLARITHROMYCIN™, and/or AZITHROMYCIN™ to prophylactically treat or prevent an opportunistic *Mycobacterium tuberculosis* infection. In another specific embodiment, Therapeutics of the invention are used in any combination with GANCICLOVIR™, FOSCARNET™, and/or CIDOFOVIR™ to prophylactically treat or prevent an opportunistic cytomegalovirus infection. In another specific embodiment, Therapeutics of the invention are used in any combination with FLUCONAZOLE™, ITRACONAZOLE™, and/or KETOCONAZOLE™ to prophylactically treat or prevent an opportunistic fungal infection. In another specific embodiment, Therapeutics of the invention are used in any combination with ACYCLOVIR™ and/or FAMCICOLVIR™ to prophylactically treat or prevent an opportunistic herpes simplex virus type I and/or type II infection. In another specific embodiment, Therapeutics of the invention are used in any combination with PYRIMETHAMINE™ and/or LEUCOVORIN™ to prophylactically treat or prevent an opportunistic *Toxoplasma gondii* infection. In another specific embodiment, Therapeutics of the invention are used in any combination with LEUCOVORIN™ and/or NEUPOGEN™ to prophylactically treat or prevent an opportunistic bacterial infection.

In a further embodiment, the Therapeutics of the invention are administered in combination with an antiviral agent. Antiviral agents that may be administered with the Therapeutics of the invention include, but are not limited to, acyclovir, ribavirin, amantadine, and remantidine.

In a further embodiment, the Therapeutics of the invention are administered in combination with an antibiotic agent. Antibiotic agents that may be administered with the Therapeutics of the invention include, but are not limited to, amoxicillin, beta-lactamases, aminoglycosides, beta-lactam (glycopeptide), beta-lactamases, Clindamycin, chloramphenicol, cephalosporins, ciprofloxacin, ciprofloxacin, erythromycin,

fluoroquinolones, macrolides, metronidazole, penicillins, quinolones, rifampin, streptomycin, sulfonamide, tetracyclines, trimethoprim, trimethoprim-sulfamthoxazole, and vancomycin.

Conventional nonspecific immunosuppressive agents, that may be administered in combination with the Therapeutics of the invention include, but are not limited to, steroids, cyclosporine, cyclosporine analogs, cyclophosphamide methylprednisone, prednisone, azathioprine, FK-506, 15-deoxyspergualin, and other immunosuppressive agents that act by suppressing the function of responding T cells.

In specific embodiments, Therapeutics of the invention are administered in combination with immunosuppressants. Immunosuppressants preparations that may be administered with the Therapeutics of the invention include, but are not limited to, ORTHOCLONE™ (OKT3), SANDIMMUNE™/NEORAL™/SANGDYA™ (cyclosporin), PROGRAF™ (tacrolimus), CELLCEPT™ (mycophenolate), Azathioprine, glucocorticosteroids, and RAPAMUNE™ (sirolimus). In a specific embodiment, immunosuppressants may be used to prevent rejection of organ or bone marrow transplantation.

In an additional embodiment, Therapeutics of the invention are administered alone or in combination with one or more intravenous immune globulin preparations. Intravenous immune globulin preparations that may be administered with the Therapeutics of the invention include, but not limited to, GAMMAR™, IVEEGAM™, SANDOGLOBULIN™, GAMMAGARD S/D™, and GAMIMUNE™. In a specific embodiment, Therapeutics of the invention are administered in combination with intravenous immune globulin preparations in transplantation therapy (e.g., bone marrow transplant).

In an additional embodiment, the Therapeutics of the invention are administered alone or in combination with an anti-inflammatory agent. Anti-inflammatory agents that may be administered with the Therapeutics of the invention include, but are not limited to, glucocorticoids and the nonsteroidal anti-inflammatories, aminoarylcarboxylic acid derivatives, arylacetic acid derivatives, arylbutyric acid derivatives, arylcarboxylic acids, arylpropionic acid derivatives, pyrazoles, pyrazolones, salicylic acid derivatives, thiazinecarboxamides, e-acetamidocaproic acid, S-adenosylmethionine, 3-amino-4-hydroxybutyric acid, amixetrine, bendazac, benzydamine, bucolome, difenpiramide, ditazol, emorfazone, guaiazulene, nabumetone, nimesulide, orgotein, oxaceprol, paranyline, perisoxal, pifoxime, proquazone, proxazole, and tenidap.

In an additional embodiment, the compositions of the invention are administered alone or in combination with an anti-angiogenic agent. Anti-angiogenic agents that may be administered with the compositions of the invention include, but are not limited to, Angiostatin (Entremed, Rockville, MD), Troponin-1 (Boston Life Sciences, Boston, MA),
5 anti-Invasive Factor, retinoic acid and derivatives thereof, paclitaxel (Taxol), Suramin, Tissue Inhibitor of Metalloproteinase-1, Tissue Inhibitor of Metalloproteinase-2, VEGI, Plasminogen Activator Inhibitor-1, Plasminogen Activator Inhibitor-2, and various forms of the lighter "d group" transition metals.

Lighter "d group" transition metals include, for example, vanadium, molybdenum,
10 tungsten, titanium, niobium, and tantalum species. Such transition metal species may form transition metal complexes. Suitable complexes of the above-mentioned transition metal species include oxo transition metal complexes.

Representative examples of vanadium complexes include oxo vanadium complexes such as vanadate and vanadyl complexes. Suitable vanadate complexes include
15 metavanadate and orthovanadate complexes such as, for example, ammonium metavanadate, sodium metavanadate, and sodium orthovanadate. Suitable vanadyl complexes include, for example, vanadyl acetylacetonate and vanadyl sulfate including vanadyl sulfate hydrates such as vanadyl sulfate mono- and trihydrates.

Representative examples of tungsten and molybdenum complexes also include oxo
20 complexes. Suitable oxo tungsten complexes include tungstate and tungsten oxide complexes. Suitable tungstate complexes include ammonium tungstate, calcium tungstate, sodium tungstate dihydrate, and tungstic acid. Suitable tungsten oxides include tungsten (IV) oxide and tungsten (VI) oxide. Suitable oxo molybdenum complexes include molybdate, molybdenum oxide, and molybdenyl complexes. Suitable molybdate complexes include
25 ammonium molybdate and its hydrates, sodium molybdate and its hydrates, and potassium molybdate and its hydrates. Suitable molybdenum oxides include molybdenum (VI) oxide, molybdenum (VI) oxide, and molybdic acid. Suitable molybdenyl complexes include, for example, molybdenyl acetylacetonate. Other suitable tungsten and molybdenum complexes include hydroxo derivatives derived from, for example, glycerol, tartaric acid, and sugars.

30 A wide variety of other anti-angiogenic factors may also be utilized within the context of the present invention. Representative examples include, but are not limited to, platelet factor 4; protamine sulphate; sulphated chitin derivatives (prepared from queen crab shells),

(Murata et al., Cancer Res. 51:22-26, 1991); Sulphated Polysaccharide Peptidoglycan Complex (SP- PG) (the function of this compound may be enhanced by the presence of steroids such as estrogen, and tamoxifen citrate); Staurosporine; modulators of matrix metabolism, including for example, proline analogs, cishydroxyproline, d,L-3,4-dehydroproline, Thiaproline, alpha,alpha-dipyridyl, aminopropionitrile fumarate; 4-propyl-5-(4-pyridinyl)-2(3H)-oxazolone; Methotrexate; Mitoxantrone; Heparin; Interferons; 2 Macroglobulin-serum; ChIMP-3 (Pavloff et al., J. Bio. Chem. 267:17321-17326, 1992); Chymostatin (Tomkinson et al., Biochem J. 286:475-480, 1992); Cyclodextrin Tetradasulfate; Eponemycin; Camptothecin; Fumagillin (Ingber et al., Nature 348:555-557, 10 1990); Gold Sodium Thiomalate ("GST"; Matsubara and Ziff, J. Clin. Invest. 79:1440-1446, 1987); anticollagenase-serum; alpha2-antiplasmin (Holmes et al., J. Biol. Chem. 262(4):1659-1664, 1987); Bisantrene (National Cancer Institute); Lobenzarit disodium (N-(2)-carboxyphenyl-4- chloroanthronilic acid disodium or "CCA"; (Takeuchi et al., Agents Actions 36:312-316, 1992); and metalloproteinase inhibitors such as BB94.

15 Additional anti-angiogenic factors that may also be utilized within the context of the present invention include Thalidomide, (Celgene, Warren, NJ); Angiostatic steroid; AGM-1470 (H. Brem and J. Folkman *J Pediatr. Surg.* 28:445-51 (1993)); an integrin alpha-v beta 3 antagonist (C. Storgard et al., *J Clin. Invest.* 103:47-54 (1999)); carboxynaminolmidazole; Carboxyamidotriazole (CAI) (National Cancer Institute, Bethesda, MD); Conbretastatin A-4 20 (CA4P) (OXiGENE, Boston, MA); Squalamine (Magainin Pharmaceuticals, Plymouth Meeting, PA); TNP-470, (Tap Pharmaceuticals, Deerfield, IL); ZD-0101 AstraZeneca (London, UK); APRA (CT2584); Benefin, Byrostatin-1 (SC339555); CGP-41251 (PKC 412); CM101; Dexrazoxane (ICRF187); DMXAA; Endostatin; Flavopridiol; Genestein; GTE; ImmTher; Iressa (ZD1839); Octreotide (Somatostatin); Panretin; Penacillamine; 25 Photopoint; PI-88; Prinomastat (AG-3340) Purlytin; Suradista (FCE26644); Tamoxifen (Nolvadex); Tazarotene; Tetrathiomolybdate; Xeloda (Capecitabine); and 5-Fluorouracil.

Anti-angiogenic agents that may be administered in combination with the compounds of the invention may work through a variety of mechanisms including, but not limited to, inhibiting proteolysis of the extracellular matrix, blocking the function of endothelial cell-extracellular matrix adhesion molecules, by antagonizing the function of angiogenesis 30 inducers such as growth factors, and inhibiting integrin receptors expressed on proliferating endothelial cells. Examples of anti-angiogenic inhibitors that interfere with extracellular

matrix proteolysis and which may be administered in combination with the compositions of the invention include, but are not limited to, AG-3340 (Agouron, La Jolla, CA), BAY-12-9566 (Bayer, West Haven, CT), BMS-275291 (Bristol Myers Squibb, Princeton, NJ), CGS-27032A (Novartis, East Hanover, NJ), Marimastat (British Biotech, Oxford, UK), and Metastat (Aeterna, St-Foy, Quebec). Examples of anti-angiogenic inhibitors that act by blocking the function of endothelial cell-extracellular matrix adhesion molecules and which may be administered in combination with the compositions of the invention include, but are not limited to, EMD-121974 (Merck KGaA Darmstadt, Germany) and Vitaxin (Ixsys, La Jolla, CA/Medimmune, Gaithersburg, MD). Examples of anti-angiogenic agents that act by directly antagonizing or inhibiting angiogenesis inducers and which may be administered in combination with the compositions of the invention include, but are not limited to, Angiozyme (Ribozyme, Boulder, CO), Anti-VEGF antibody (Genentech, S. San Francisco, CA), PTK-787/ZK-225846 (Novartis, Basel, Switzerland), SU-101 (Sugen, S. San Francisco, CA), SU-5416 (Sugen/ Pharmacia Upjohn, Bridgewater, NJ), and SU-6668 (Sugen). Other anti-angiogenic agents act to indirectly inhibit angiogenesis. Examples of indirect inhibitors of angiogenesis which may be administered in combination with the compositions of the invention include, but are not limited to, IM-862 (Cytran, Kirkland, WA), Interferon-alpha, IL-12 (Roche, Nutley, NJ), and Pentosan polysulfate (Georgetown University, Washington, DC).

In particular embodiments, the use of compositions of the invention in combination with anti-angiogenic agents is contemplated for the treatment, prevention, and/or amelioration of an autoimmune disease, such as for example, an autoimmune disease described herein.

In a particular embodiment, the use of compositions of the invention in combination with anti-angiogenic agents is contemplated for the treatment, prevention, and/or amelioration of arthritis. In a more particular embodiment, the use of compositions of the invention in combination with anti-angiogenic agents is contemplated for the treatment, prevention, and/or amelioration of rheumatoid arthritis.

In another embodiment, compositions of the invention are administered in combination with a chemotherapeutic agent. Chemotherapeutic agents that may be administered with the Therapeutics of the invention include, but are not limited to, antibiotic derivatives (e.g., doxorubicin, bleomycin, daunorubicin, and dactinomycin); antiestrogens

(e.g., tamoxifen); antimetabolites (e.g., fluorouracil, 5-FU, methotrexate, floxuridine, interferon alpha-2b, glutamic acid, plicamycin, mercaptopurine, and 6-thioguanine); cytotoxic agents (e.g., carmustine, BCNU, lomustine, CCNU, cytosine arabinoside, cyclophosphamide, estramustine, hydroxyurea, procarbazine, mitomycin, busulfan, cis-platin, and vincristine sulfate); hormones (e.g., medroxyprogesterone, estramustine phosphate sodium, ethinyl estradiol, estradiol, megestrol acetate, methyltestosterone, diethylstilbestrol diphosphate, chlorotrianisene, and testolactone); nitrogen mustard derivatives (e.g., mephallen, chorambucil, mechlorethamine (nitrogen mustard) and thiotepa); steroids and combinations (e.g., bethamethasone sodium phosphate); and others (e.g., dicarbazine, asparaginase, mitotane, vincristine sulfate, vinblastine sulfate, and etoposide).

In a specific embodiment, Therapeutics of the invention are administered in combination with CHOP (cyclophosphamide, doxorubicin, vincristine, and prednisone) or any combination of the components of CHOP. In another embodiment, Therapeutics of the invention are administered in combination with Rituximab. In a further embodiment, Therapeutics of the invention are administered with Rituxmab and CHOP, or Rituxmab and any combination of the components of CHOP.

In an additional embodiment, the Therapeutics of the invention are administered in combination with cytokines. Cytokines that may be administered with the Therapeutics of the invention include, but are not limited to, IL2, IL3, IL4, IL5, IL6, IL7, IL10, IL12, IL13, IL15, anti-CD40, CD40L, IFN-gamma and TNF-alpha. In another embodiment, Therapeutics of the invention may be administered with any interleukin, including, but not limited to, IL-1alpha, IL-1beta, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-11, IL-12, IL-13, IL-14, IL-15, IL-16, IL-17, IL-18, IL-19, IL-20, and IL-21.

In an additional embodiment, the Therapeutics of the invention are administered in combination with angiogenic proteins. Angiogenic proteins that may be administered with the Therapeutics of the invention include, but are not limited to, Glioma Derived Growth Factor (GDGF), as disclosed in European Patent Number EP-399816; Platelet Derived Growth Factor-A (PDGF-A), as disclosed in European Patent Number EP-682110; Platelet Derived Growth Factor-B (PDGF-B), as disclosed in European Patent Number EP-282317; Placental Growth Factor (PIGF), as disclosed in International Publication Number WO 92/06194; Placental Growth Factor-2 (PIGF-2), as disclosed in Hauser et al., Growth Factors, 4:259-268 (1993); Vascular Endothelial Growth Factor (VEGF), as disclosed in International

Publication Number WO 90/13649; Vascular Endothelial Growth Factor-A (VEGF-A), as disclosed in European Patent Number EP-506477; Vascular Endothelial Growth Factor-2 (VEGF-2), as disclosed in International Publication Number WO 96/39515; Vascular Endothelial Growth Factor B (VEGF-3); Vascular Endothelial Growth Factor B-186 (VEGF-B186), as disclosed in International Publication Number WO 96/26736; Vascular Endothelial Growth Factor-D (VEGF-D), as disclosed in International Publication Number WO 98/02543; Vascular Endothelial Growth Factor-D (VEGF-D), as disclosed in International Publication Number WO 98/07832; and Vascular Endothelial Growth Factor-E (VEGF-E), as disclosed in German Patent Number DE19639601. The above mentioned references are incorporated herein by reference herein.

In an additional embodiment, the Therapeutics of the invention are administered in combination with hematopoietic growth factors. Hematopoietic growth factors that may be administered with the Therapeutics of the invention include, but are not limited to, LEUKINE™ (SARGRAMOSTIM™) and NEUPOGEN™ (FILGRASTIM™).

In an additional embodiment, the Therapeutics of the invention are administered in combination with Fibroblast Growth Factors. Fibroblast Growth Factors that may be administered with the Therapeutics of the invention include, but are not limited to, FGF-1, FGF-2, FGF-3, FGF-4, FGF-5, FGF-6, FGF-7, FGF-8, FGF-9, FGF-10, FGF-11, FGF-12, FGF-13, FGF-14, and FGF-15.

In additional embodiments, the Therapeutics of the invention are administered in combination with other therapeutic or prophylactic regimens, such as, for example, radiation therapy.

Example 14: Method of Treating Decreased Levels of the Polypeptide

The present invention relates to a method for treating an individual in need of an increased level of a polypeptide of the invention in the body comprising administering to such an individual a composition comprising a therapeutically effective amount of an agonist of the invention (including polypeptides of the invention). Moreover, it will be appreciated that conditions caused by a decrease in the standard or normal expression level of a polypeptide of the present invention in an individual can be treated by administering the agonist or antagonist of the present invention. Thus, the invention also provides a method of treatment of an individual in need of an increased level of the polypeptide comprising

administering to such an individual a Therapeutic comprising an amount of the agonist or antagonist to increase the activity level of the polypeptide in such an individual.

For example, a patient with decreased levels of a polypeptide receives a daily dose 0.1-100 ug/kg of the agonist or antagonist for six consecutive days. The exact details of the dosing scheme, based on administration and formulation, are provided in Example 13.

Example 15: Method of Treating Increased Levels of the Polypeptide

The present invention also relates to a method of treating an individual in need of a decreased level of a polypeptide of the invention in the body comprising administering to such an individual a composition comprising a therapeutically effective amount of an antagonist of the invention (including polypeptides and antibodies of the invention).

In one example, antisense technology is used to inhibit production of a polypeptide of the present invention. This technology is one example of a method of decreasing levels of a polypeptide, due to a variety of etiologies, such as cancer.

For example, a patient diagnosed with abnormally increased levels of a polypeptide is administered intravenously antisense polynucleotides at 0.5, 1.0, 1.5, 2.0 and 3.0 mg/kg day for 21 days. This treatment is repeated after a 7-day rest period if the treatment was well tolerated. The formulation of the antisense polynucleotide is provided in Example 13.

Example 16: Method of Treatment Using Gene Therapy-Ex Vivo

One method of gene therapy transplants fibroblasts, which are capable of expressing a polypeptide, onto a patient. Generally, fibroblasts are obtained from a subject by skin biopsy. The resulting tissue is placed in tissue-culture medium and separated into small pieces. Small chunks of the tissue are placed on a wet surface of a tissue culture flask, approximately ten pieces are placed in each flask. The flask is turned upside down, closed tight and left at room temperature over night. After 24 hours at room temperature, the flask is inverted and the chunks of tissue remain fixed to the bottom of the flask and fresh media (e.g., Ham's F12 media, with 10% FBS, penicillin and streptomycin) is added. The flasks are then incubated at 37 degree C for approximately one week.

At this time, fresh media is added and subsequently changed every several days. After an additional two weeks in culture, a monolayer of fibroblasts emerge. The monolayer is trypsinized and scaled into larger flasks.

pMV-7 (Kirschmeier, P.T. et al., DNA, 7:219-25 (1988)), flanked by the long terminal repeats of the Moloney murine sarcoma virus, is digested with EcoRI and HindIII and subsequently treated with calf intestinal phosphatase. The linear vector is fractionated on agarose gel and purified, using glass beads.

5 The cDNA encoding a polypeptide of the present invention can be amplified using PCR primers which correspond to the 5' and 3' end sequences respectively as set forth in Example 1 using primers and having appropriate restriction sites and initiation/stop codons, if necessary. Preferably, the 5' primer contains an EcoRI site and the 3' primer includes a HindIII site. Equal quantities of the Moloney murine sarcoma virus linear backbone and the
10 amplified EcoRI and HindIII fragment are added together, in the presence of T4 DNA ligase. The resulting mixture is maintained under conditions appropriate for ligation of the two fragments. The ligation mixture is then used to transform bacteria HB101, which are then plated onto agar containing kanamycin for the purpose of confirming that the vector has the gene of interest properly inserted.

15 The amphotropic pA317 or GP+am12 packaging cells are grown in tissue culture to confluent density in Dulbecco's Modified Eagles Medium (DMEM) with 10% calf serum (CS), penicillin and streptomycin. The MSV vector containing the gene is then added to the media and the packaging cells transduced with the vector. The packaging cells now produce infectious viral particles containing the gene (the packaging cells are now referred to as
20 producer cells).

Fresh media is added to the transduced producer cells, and subsequently, the media is harvested from a 10 cm plate of confluent producer cells. The spent media, containing the infectious viral particles, is filtered through a millipore filter to remove detached producer cells and this media is then used to infect fibroblast cells. Media is removed from a sub-
25 confluent plate of fibroblasts and quickly replaced with the media from the producer cells. This media is removed and replaced with fresh media. If the titer of virus is high, then virtually all fibroblasts will be infected and no selection is required. If the titer is very low, then it is necessary to use a retroviral vector that has a selectable marker, such as neo or his. Once the fibroblasts have been efficiently infected, the fibroblasts are analyzed to determine
30 whether protein is produced.

The engineered fibroblasts are then transplanted onto the host, either alone or after having been grown to confluence on cytodex 3 microcarrier beads.

Example 17: Gene Therapy Using Endogenous Genes Corresponding To Polynucleotides of the Invention

Another method of gene therapy according to the present invention involves operably associating the endogenous polynucleotide sequence of the invention with a promoter via homologous recombination as described, for example, in U.S. Patent NO: 5,641,670, issued June 24, 1997; International Publication NO: WO 96/29411, published September 26, 1996; International Publication NO: WO 94/12650, published August 4, 1994; Koller et al., *Proc. Natl. Acad. Sci. USA*, 86:8932-8935 (1989); and Zijlstra et al., *Nature*, 342:435-438 (1989).

This method involves the activation of a gene which is present in the target cells, but which is not expressed in the cells, or is expressed at a lower level than desired.

Polynucleotide constructs are made which contain a promoter and targeting sequences, which are homologous to the 5' non-coding sequence of endogenous polynucleotide sequence, flanking the promoter. The targeting sequence will be sufficiently near the 5' end of the polynucleotide sequence so the promoter will be operably linked to the endogenous sequence upon homologous recombination. The promoter and the targeting sequences can be amplified using PCR. Preferably, the amplified promoter contains distinct restriction enzyme sites on the 5' and 3' ends. Preferably, the 3' end of the first targeting sequence contains the same restriction enzyme site as the 5' end of the amplified promoter and the 5' end of the second targeting sequence contains the same restriction site as the 3' end of the amplified promoter.

The amplified promoter and the amplified targeting sequences are digested with the appropriate restriction enzymes and subsequently treated with calf intestinal phosphatase. The digested promoter and digested targeting sequences are added together in the presence of T4 DNA ligase. The resulting mixture is maintained under conditions appropriate for ligation of the two fragments. The construct is size fractionated on an agarose gel then purified by phenol extraction and ethanol precipitation.

In this Example, the polynucleotide constructs are administered as naked polynucleotides via electroporation. However, the polynucleotide constructs may also be administered with transfection-facilitating agents, such as liposomes, viral sequences, viral particles, precipitating agents, etc. Such methods of delivery are known in the art.

Once the cells are transfected, homologous recombination will take place which results in the promoter being operably linked to the endogenous polynucleotide sequence. This results in the expression of polynucleotide corresponding to the polynucleotide in the cell. Expression may be detected by immunological staining, or any other method known in the art.

Fibroblasts are obtained from a subject by skin biopsy. The resulting tissue is placed in DMEM + 10% fetal calf serum. Exponentially growing or early stationary phase fibroblasts are trypsinized and rinsed from the plastic surface with nutrient medium. An aliquot of the cell suspension is removed for counting, and the remaining cells are subjected to centrifugation. The supernatant is aspirated and the pellet is resuspended in 5 ml of electroporation buffer (20 mM HEPES pH 7.3, 137 mM NaCl, 5 mM KCl, 0.7 mM Na₂HPO₄, 6 mM dextrose). The cells are recentrifuged, the supernatant aspirated, and the cells resuspended in electroporation buffer containing 1 mg/ml acetylated bovine serum albumin. The final cell suspension contains approximately 3×10^6 cells/ml. Electroporation should be performed immediately following resuspension.

Plasmid DNA is prepared according to standard techniques. For example, to construct a plasmid for targeting to the locus corresponding to the polynucleotide of the invention, plasmid pUC18 (MBI Fermentas, Amherst, NY) is digested with HindIII. The CMV promoter is amplified by PCR with an XbaI site on the 5' end and a BamHI site on the 3' end. Two non-coding sequences are amplified via PCR: one non-coding sequence (fragment 1) is amplified with a HindIII site at the 5' end and an Xba site at the 3' end; the other non-coding sequence (fragment 2) is amplified with a BamHI site at the 5' end and a HindIII site at the 3' end. The CMV promoter and the fragments (1 and 2) are digested with the appropriate enzymes (CMV promoter - XbaI and BamHI; fragment 1 - XbaI; fragment 2 - BamHI) and ligated together. The resulting ligation product is digested with HindIII, and ligated with the HindIII-digested pUC18 plasmid.

Plasmid DNA is added to a sterile cuvette with a 0.4 cm electrode gap (Bio-Rad). The final DNA concentration is generally at least 120 µg/ml. 0.5 ml of the cell suspension (containing approximately 1.5×10^6 cells) is then added to the cuvette, and the cell suspension and DNA solutions are gently mixed. Electroporation is performed with a Gene-Pulser apparatus (Bio-Rad). Capacitance and voltage are set at 960 µF and 250-300 V, respectively. As voltage increases, cell survival decreases, but the percentage of surviving

cells that stably incorporate the introduced DNA into their genome increases dramatically. Given these parameters, a pulse time of approximately 14-20 mSec should be observed.

Electroporated cells are maintained at room temperature for approximately 5 min, and the contents of the cuvette are then gently removed with a sterile transfer pipette. The cells are added directly to 10 ml of prewarmed nutrient media (DMEM with 15% calf serum) in a 10 cm dish and incubated at 37 degree C. The following day, the media is aspirated and replaced with 10 ml of fresh media and incubated for a further 16-24 hours.

The engineered fibroblasts are then injected into the host, either alone or after having been grown to confluence on cytodex 3 microcarrier beads. The fibroblasts now produce the protein product. The fibroblasts can then be introduced into a patient as described above.

Example 18: Method of Treatment Using Gene Therapy - In Vivo

Another aspect of the present invention is using *in vivo* gene therapy methods to treat disorders, diseases and conditions. The gene therapy method relates to the introduction of naked nucleic acid (DNA, RNA, and antisense DNA or RNA) sequences into an animal to increase or decrease the expression of the polypeptide. The polynucleotide of the present invention may be operatively linked to a promoter or any other genetic elements necessary for the expression of the polypeptide by the target tissue. Such gene therapy and delivery techniques and methods are known in the art, see, for example, WO90/11092, WO98/11779; U.S. Patent NO. 5693622, 5705151, 5580859; Tabata et al., Cardiovasc. Res. 35(3):470-479 (1997); Chao et al., Pharmacol. Res. 35(6):517-522 (1997); Wolff, Neuromuscul. Disord. 7(5):314-318 (1997); Schwartz et al., Gene Ther. 3(5):405-411 (1996); Tsurumi et al., Circulation 94(12):3281-3290 (1996) (incorporated herein by reference).

The polynucleotide constructs may be delivered by any method that delivers injectable materials to the cells of an animal, such as, injection into the interstitial space of tissues (heart, muscle, skin, lung, liver, intestine and the like). The polynucleotide constructs can be delivered in a pharmaceutically acceptable liquid or aqueous carrier.

The term "naked" polynucleotide, DNA or RNA, refers to sequences that are free from any delivery vehicle that acts to assist, promote, or facilitate entry into the cell, including viral sequences, viral particles, liposome formulations, lipofectin or precipitating agents and the like. However, the polynucleotides of the present invention may also be delivered in liposome formulations (such as those taught in Felgner P.L. et al. (1995) Ann.

NY Acad. Sci. 772:126-139 and Abdallah B. et al. (1995) Biol. Cell 85(1):1-7) which can be prepared by methods well known to those skilled in the art.

The polynucleotide vector constructs used in the gene therapy method are preferably constructs that will not integrate into the host genome nor will they contain sequences that allow for replication. Any strong promoter known to those skilled in the art can be used for driving the expression of DNA. Unlike other gene therapies techniques, one major advantage of introducing naked nucleic acid sequences into target cells is the transitory nature of the polynucleotide synthesis in the cells. Studies have shown that non-replicating DNA sequences can be introduced into cells to provide production of the desired polypeptide for periods of up to six months.

The polynucleotide construct can be delivered to the interstitial space of tissues within the an animal, including of muscle, skin, brain, lung, liver, spleen, bone marrow, thymus, heart, lymph, blood, bone, cartilage, pancreas, kidney, gall bladder, stomach, intestine, testis, ovary, uterus, rectum, nervous system, eye, gland, and connective tissue. Interstitial space of the tissues comprises the intercellular fluid, mucopolysaccharide matrix among the reticular fibers of organ tissues, elastic fibers in the walls of vessels or chambers, collagen fibers of fibrous tissues, or that same matrix within connective tissue ensheathing muscle cells or in the lacunae of bone. It is similarly the space occupied by the plasma of the circulation and the lymph fluid of the lymphatic channels. Delivery to the interstitial space of muscle tissue is preferred for the reasons discussed below. They may be conveniently delivered by injection into the tissues comprising these cells. They are preferably delivered to and expressed in persistent, non-dividing cells which are differentiated, although delivery and expression may be achieved in non-differentiated or less completely differentiated cells, such as, for example, stem cells of blood or skin fibroblasts. *In vivo* muscle cells are particularly competent in their ability to take up and express polynucleotides.

For the naked polynucleotide injection, an effective dosage amount of DNA or RNA will be in the range of from about 0.05 g/kg body weight to about 50 mg/kg body weight. Preferably the dosage will be from about 0.005 mg/kg to about 20 mg/kg and more preferably from about 0.05 mg/kg to about 5 mg/kg. Of course, as the artisan of ordinary skill will appreciate, this dosage will vary according to the tissue site of injection. The appropriate and effective dosage of nucleic acid sequence can readily be determined by those of ordinary skill in the art and may depend on the condition being treated and the route of administration. The

preferred route of administration is by the parenteral route of injection into the interstitial space of tissues. However, other parenteral routes may also be used, such as, inhalation of an aerosol formulation particularly for delivery to lungs or bronchial tissues, throat or mucous membranes of the nose. In addition, naked polynucleotide constructs can be delivered to
5 arteries during angioplasty by the catheter used in the procedure.

The dose response effects of injected polynucleotide in muscle *in vivo* is determined as follows. Suitable template DNA for production of mRNA coding for polypeptide of the present invention is prepared in accordance with a standard recombinant DNA methodology. The template DNA, which may be either circular or linear, is either used as naked DNA or
10 complexed with liposomes. The quadriceps muscles of mice are then injected with various amounts of the template DNA.

Five to six week old female and male Balb/C mice are anesthetized by intraperitoneal injection with 0.3 ml of 2.5% Avertin. A 1.5 cm incision is made on the anterior thigh, and the quadriceps muscle is directly visualized. The template DNA is injected in 0.1 ml of
15 carrier in a 1 cc syringe through a 27 gauge needle over one minute, approximately 0.5 cm from the distal insertion site of the muscle into the knee and about 0.2 cm deep. A suture is placed over the injection site for future localization, and the skin is closed with stainless steel clips.

After an appropriate incubation time (e.g., 7 days) muscle extracts are prepared by
20 excising the entire quadriceps. Every fifth 15 um cross-section of the individual quadriceps muscles is histochemically stained for protein expression. A time course for protein expression may be done in a similar fashion except that quadriceps from different mice are harvested at different times. Persistence of DNA in muscle following injection may be determined by Southern blot analysis after preparing total cellular DNA and HIRT
25 supernatants from injected and control mice. The results of the above experimentation in mice can be use to extrapolate proper dosages and other treatment parameters in humans and other animals using naked DNA.

Example 19: Transgenic Animals

30 The polypeptides of the invention can also be expressed in transgenic animals. Animals of any species, including, but not limited to, mice, rats, rabbits, hamsters, guinea pigs, pigs, micro-pigs, goats, sheep, cows and non-human primates, e.g., baboons, monkeys,

and chimpanzees may be used to generate transgenic animals. In a specific embodiment, techniques described herein or otherwise known in the art, are used to express polypeptides of the invention in humans, as part of a gene therapy protocol.

Any technique known in the art may be used to introduce the transgene (i.e., polynucleotides of the invention) into animals to produce the founder lines of transgenic animals. Such techniques include, but are not limited to, pronuclear microinjection (Paterson et al., Appl. Microbiol. Biotechnol. 40:691-698 (1994); Carver et al., Biotechnology (NY) 11:1263-1270 (1993); Wright et al., Biotechnology (NY) 9:830-834 (1991); and Hoppe et al., U.S. Pat. No. 4,873,191 (1989)); retrovirus mediated gene transfer into germ lines (Van der Putten et al., Proc. Natl. Acad. Sci., USA 82:6148-6152 (1985)), blastocysts or embryos; gene targeting in embryonic stem cells (Thompson et al., Cell 56:313-321 (1989)); electroporation of cells or embryos (Lo, 1983, Mol Cell. Biol. 3:1803-1814 (1983)); introduction of the polynucleotides of the invention using a gene gun (see, e.g., Ulmer et al., Science 259:1745 (1993); introducing nucleic acid constructs into embryonic pluripotent stem cells and transferring the stem cells back into the blastocyst; and sperm-mediated gene transfer (Lavitrano et al., Cell 57:717-723 (1989); etc. For a review of such techniques, see Gordon, "Transgenic Animals," Intl. Rev. Cytol. 115:171-229 (1989), which is incorporated by reference herein in its entirety.

Any technique known in the art may be used to produce transgenic clones containing polynucleotides of the invention, for example, nuclear transfer into enucleated oocytes of nuclei from cultured embryonic, fetal, or adult cells induced to quiescence (Campbell et al., Nature 380:64-66 (1996); Wilmut et al., Nature 385:810-813 (1997)).

The present invention provides for transgenic animals that carry the transgene in all their cells, as well as animals which carry the transgene in some, but not all their cells, i.e., mosaic animals or chimeric. The transgene may be integrated as a single transgene or as multiple copies such as in concatamers, e.g., head-to-head tandems or head-to-tail tandems. The transgene may also be selectively introduced into and activated in a particular cell type by following, for example, the teaching of Lasko et al. (Lasko et al., Proc. Natl. Acad. Sci. USA 89:6232-6236 (1992)). The regulatory sequences required for such a cell-type specific activation will depend upon the particular cell type of interest, and will be apparent to those of skill in the art. When it is desired that the polynucleotide transgene be integrated into the chromosomal site of the endogenous gene, gene targeting is preferred. Briefly, when such a

technique is to be utilized, vectors containing some nucleotide sequences homologous to the endogenous gene are designed for the purpose of integrating, via homologous recombination with chromosomal sequences, into and disrupting the function of the nucleotide sequence of the endogenous gene. The transgene may also be selectively introduced into a particular cell type, thus inactivating the endogenous gene in only that cell type, by following, for example, the teaching of Gu et al. (Gu et al., Science 265:103-106 (1994)). The regulatory sequences required for such a cell-type specific inactivation will depend upon the particular cell type of interest, and will be apparent to those of skill in the art.

Once transgenic animals have been generated, the expression of the recombinant gene may be assayed utilizing standard techniques. Initial screening may be accomplished by Southern blot analysis or PCR techniques to analyze animal tissues to verify that integration of the transgene has taken place. The level of mRNA expression of the transgene in the tissues of the transgenic animals may also be assessed using techniques which include, but are not limited to, Northern blot analysis of tissue samples obtained from the animal, *in situ* hybridization analysis, and reverse transcriptase-PCR (rt-PCR). Samples of transgenic gene-expressing tissue may also be evaluated immunocytochemically or immunohistochemically using antibodies specific for the transgene product.

Once the founder animals are produced, they may be bred, inbred, outbred, or crossbred to produce colonies of the particular animal. Examples of such breeding strategies include, but are not limited to: outbreeding of founder animals with more than one integration site in order to establish separate lines; inbreeding of separate lines in order to produce compound transgenics that express the transgene at higher levels because of the effects of additive expression of each transgene; crossing of heterozygous transgenic animals to produce animals homozygous for a given integration site in order to both augment expression and eliminate the need for screening of animals by DNA analysis; crossing of separate homozygous lines to produce compound heterozygous or homozygous lines; and breeding to place the transgene on a distinct background that is appropriate for an experimental model of interest.

Transgenic animals of the invention have uses which include, but are not limited to, animal model systems useful in elaborating the biological function of polypeptides of the present invention, studying conditions and/or disorders associated with aberrant expression, and in screening for compounds effective in ameliorating such conditions and/or disorders.

Example 20: Knock-Out Animals

Endogenous gene expression can also be reduced by inactivating or "knocking out" the gene and/or its promoter using targeted homologous recombination. (E.g., see Smithies et al., Nature 317:230-234 (1985); Thomas & Capecchi, Cell 51:503-512 (1987); Thompson et al., Cell 5:313-321 (1989); each of which is incorporated by reference herein in its entirety). For example, a mutant, non-functional polynucleotide of the invention (or a completely unrelated DNA sequence) flanked by DNA homologous to the endogenous polynucleotide sequence (either the coding regions or regulatory regions of the gene) can be used, with or without a selectable marker and/or a negative selectable marker, to transfect cells that express polypeptides of the invention in vivo. In another embodiment, techniques known in the art are used to generate knockouts in cells that contain, but do not express the gene of interest. Insertion of the DNA construct, via targeted homologous recombination, results in inactivation of the targeted gene. Such approaches are particularly suited in research and agricultural fields where modifications to embryonic stem cells can be used to generate animal offspring with an inactive targeted gene (e.g., see Thomas & Capecchi 1987 and Thompson 1989, *supra*). However this approach can be routinely adapted for use in humans provided the recombinant DNA constructs are directly administered or targeted to the required site in vivo using appropriate viral vectors that will be apparent to those of skill in the art.

In further embodiments of the invention, cells that are genetically engineered to express the polypeptides of the invention, or alternatively, that are genetically engineered not to express the polypeptides of the invention (e.g., knockouts) are administered to a patient in vivo. Such cells may be obtained from the patient (i.e., animal, including human) or an MHC compatible donor and can include, but are not limited to fibroblasts, bone marrow cells, blood cells (e.g., lymphocytes), adipocytes, muscle cells, endothelial cells etc. The cells are genetically engineered *in vitro* using recombinant DNA techniques to introduce the coding sequence of polypeptides of the invention into the cells, or alternatively, to disrupt the coding sequence and/or endogenous regulatory sequence associated with the polypeptides of the invention, e.g., by transduction (using viral vectors, and preferably vectors that integrate the transgene into the cell genome) or transfection procedures, including, but not limited to, the use of plasmids, cosmids, YACs, naked DNA, electroporation, liposomes, etc. The coding

sequence of the polypeptides of the invention can be placed under the control of a strong constitutive or inducible promoter or promoter/enhancer to achieve expression, and preferably secretion, of the polypeptides of the invention. The engineered cells which express and preferably secrete the polypeptides of the invention can be introduced into the patient systemically, e.g., in the circulation, or intraperitoneally.

Alternatively, the cells can be incorporated into a matrix and implanted in the body, e.g., genetically engineered fibroblasts can be implanted as part of a skin graft; genetically engineered endothelial cells can be implanted as part of a lymphatic or vascular graft. (See, for example, Anderson et al. U.S. Patent No. 5,399,349; and Mulligan & Wilson, U.S. Patent No. 5,460,959 each of which is incorporated by reference herein in its entirety).

When the cells to be administered are non-autologous or non-MHC compatible cells, they can be administered using well known techniques which prevent the development of a host immune response against the introduced cells. For example, the cells may be introduced in an encapsulated form which, while allowing for an exchange of components with the immediate extracellular environment, does not allow the introduced cells to be recognized by the host immune system.

Transgenic and "knock-out" animals of the invention have uses which include, but are not limited to, animal model systems useful in elaborating the biological function of polypeptides of the present invention, studying conditions and/or disorders associated with aberrant expression, and in screening for compounds effective in ameliorating such conditions and/or disorders.

Example 22: Assays Detecting Stimulation or Inhibition of B cell Proliferation and Differentiation

Generation of functional humoral immune responses requires both soluble and cognate signaling between B-lineage cells and their microenvironment. Signals may impart a positive stimulus that allows a B-lineage cell to continue its programmed development, or a negative stimulus that instructs the cell to arrest its current developmental pathway. To date, numerous stimulatory and inhibitory signals have been found to influence B cell responsiveness including IL-2, IL-4, IL-5, IL-6, IL-7, IL10, IL-13, IL-14 and IL-15. Interestingly, these signals are by themselves weak effectors but can, in combination with

various co-stimulatory proteins, induce activation, proliferation, differentiation, homing, tolerance and death among B cell populations.

One of the best studied classes of B-cell co-stimulatory proteins is the TNF-superfamily. Within this family CD40, CD27, and CD30 along with their respective ligands CD154, CD70, and CD153 have been found to regulate a variety of immune responses. Assays which allow for the detection and/or observation of the proliferation and differentiation of these B-cell populations and their precursors are valuable tools in determining the effects various proteins may have on these B-cell populations in terms of proliferation and differentiation. Listed below are two assays designed to allow for the detection of the differentiation, proliferation, or inhibition of B-cell populations and their precursors.

In Vitro Assay- Agonists or antagonists of the invention can be assessed for its ability to induce activation, proliferation, differentiation or inhibition and/or death in B-cell populations and their precursors. The activity of the agonists or antagonists of the invention on purified human tonsillar B cells, measured qualitatively over the dose range from 0.1 to 10,000 ng/mL, is assessed in a standard B-lymphocyte co-stimulation assay in which purified tonsillar B cells are cultured in the presence of either formalin-fixed *Staphylococcus aureus* Cowan I (SAC) or immobilized anti-human IgM antibody as the priming agent. Second signals such as IL-2 and IL-15 synergize with SAC and IgM crosslinking to elicit B cell proliferation as measured by tritiated-thymidine incorporation. Novel synergizing agents can be readily identified using this assay. The assay involves isolating human tonsillar B cells by magnetic bead (MACS) depletion of CD3-positive cells. The resulting cell population is greater than 95% B cells as assessed by expression of CD45R(B220).

Various dilutions of each sample are placed into individual wells of a 96-well plate to which are added 10^5 B-cells suspended in culture medium (RPMI 1640 containing 10% FBS, 5×10^{-5} M 2ME, 100U/ml penicillin, 10ug/ml streptomycin, and 10^{-5} dilution of SAC) in a total volume of 150ul. Proliferation or inhibition is quantitated by a 20h pulse (1uCi/well) with 3 H-thymidine (6.7 Ci/mM) beginning 72h post factor addition. The positive and negative controls are IL2 and medium respectively.

In Vivo Assay- BALB/c mice are injected (i.p.) twice per day with buffer only, or 2 mg/Kg of agonists or antagonists of the invention, or truncated forms thereof. Mice receive this treatment for 4 consecutive days, at which time they are sacrificed and various tissues

and serum collected for analyses. Comparison of H&E sections from normal spleens and spleens treated with agonists or antagonists of the invention identify the results of the activity of the agonists or antagonists on spleen cells, such as the diffusion of peri-arterial lymphatic sheaths, and/or significant increases in the nucleated cellularity of the red pulp regions, which may indicate the activation of the differentiation and proliferation of B-cell populations. Immunohistochemical studies using a B cell marker, anti-CD45R(B220), are used to determine whether any physiological changes to splenic cells, such as splenic disorganization, are due to increased B-cell representation within loosely defined B-cell zones that infiltrate established T-cell regions.

Flow cytometric analyses of the spleens from mice treated with agonist or antagonist is used to indicate whether the agonists or antagonists specifically increases the proportion of ThB+, CD45R(B220)dull B cells over that which is observed in control mice.

Likewise, a predicted consequence of increased mature B-cell representation in vivo is a relative increase in serum Ig titers. Accordingly, serum IgM and IgA levels are compared between buffer and agonists or antagonists-treated mice.

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

Example 23: T Cell Proliferation Assay

A CD3-induced proliferation assay is performed on PBMCs and is measured by the uptake of ^3H -thymidine. The assay is performed as follows. Ninety-six well plates are coated with 100 μl /well of mAb to CD3 (HIT3a, Pharmingen) or isotype-matched control mAb (B33.1) overnight at 4 degrees C (1 $\mu\text{g}/\text{ml}$ in .05M bicarbonate buffer, pH 9.5), then washed three times with PBS. PBMC are isolated by F/H gradient centrifugation from human peripheral blood and added to quadruplicate wells (5×10^4 /well) of mAb coated plates in RPMI containing 10% FCS and P/S in the presence of varying concentrations of agonists or antagonists of the invention (total volume 200. μl). Relevant protein buffer and medium alone are controls. After 48 hr. culture at 37 degrees C, plates are spun for 2 min. at 1000 rpm and 100 μl of supernatant is removed and stored -20 degrees C for measurement of IL-2 (or other cytokines) if effect on proliferation is observed. Wells are supplemented with 100 μl of medium containing 0.5 uCi of ^3H -thymidine and cultured at 37 degrees C for 18-24 hr.

Wells are harvested and incorporation of ^3H -thymidine used as a measure of proliferation. Anti-CD3 alone is the positive control for proliferation. IL-2 (100 U/ml) is also used as a control which enhances proliferation. Control antibody which does not induce proliferation of T cells is used as the negative controls for the effects of agonists or antagonists of the invention.

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

Example 24: Effect of Agonists or Antagonists of the Invention on the Expression of MHC Class II, Costimulatory and Adhesion Molecules and Cell Differentiation of Monocytes and Monocyte-Derived Human Dendritic Cells

Dendritic cells are generated by the expansion of proliferating precursors found in the peripheral blood: adherent PBMC or elutriated monocytic fractions are cultured for 7-10 days with GM-CSF (50 ng/ml) and IL-4 (20 ng/ml). These dendritic cells have the characteristic phenotype of immature cells (expression of CD1, CD80, CD86, CD40 and MHC class II antigens). Treatment with activating factors, such as TNF- α , causes a rapid change in surface phenotype (increased expression of MHC class I and II, costimulatory and adhesion molecules, downregulation of FC γ RII, upregulation of CD83). These changes correlate with increased antigen-presenting capacity and with functional maturation of the dendritic cells.

FACS analysis of surface antigens is performed as follows. Cells are treated 1-3 days with increasing concentrations of agonist or antagonist of the invention or LPS (positive control), washed with PBS containing 1% BSA and 0.02 mM sodium azide, and then incubated with 1:20 dilution of appropriate FITC- or PE-labeled monoclonal antibodies for 30 minutes at 4 degrees C. After an additional wash, the labeled cells are analyzed by flow cytometry on a FACScan (Becton Dickinson).

Effect on the production of cytokines. Cytokines generated by dendritic cells, in particular IL-12, are important in the initiation of T-cell dependent immune responses. IL-12 strongly influences the development of Th1 helper T-cell immune response, and induces cytotoxic T and NK cell function. An ELISA is used to measure the IL-12 release as follows.

Dendritic cells ($10^6/\text{ml}$) are treated with increasing concentrations of agonists or antagonists of the invention for 24 hours. LPS (100 ng/ml) is added to the cell culture as positive control. Supernatants from the cell cultures are then collected and analyzed for IL-12 content using commercial ELISA kit (e.g., R & D Systems (Minneapolis, MN)). The standard protocols provided with the kits are used.

Effect on the expression of MHC Class II, costimulatory and adhesion molecules.

Three major families of cell surface antigens can be identified on monocytes: adhesion molecules, molecules involved in antigen presentation, and Fc receptor. Modulation of the expression of MHC class II antigens and other costimulatory molecules, such as B7 and ICAM-1, may result in changes in the antigen presenting capacity of monocytes and ability to induce T cell activation. Increase expression of Fc receptors may correlate with improved monocyte cytotoxic activity, cytokine release and phagocytosis.

FACS analysis is used to examine the surface antigens as follows. Monocytes are treated 1-5 days with increasing concentrations of agonists or antagonists of the invention or LPS (positive control), washed with PBS containing 1% BSA and 0.02 mM sodium azide, and then incubated with 1:20 dilution of appropriate FITC- or PE-labeled monoclonal antibodies for 30 minutes at 4 degreesC. After an additional wash, the labeled cells are analyzed by flow cytometry on a FACScan (Becton Dickinson).

Monocyte activation and/or increased survival. Assays for molecules that activate (or alternatively, inactivate) monocytes and/or increase monocyte survival (or alternatively, decrease monocyte survival) are known in the art and may routinely be applied to determine whether a molecule of the invention functions as an inhibitor or activator of monocytes. Agonists or antagonists of the invention can be screened using the three assays described below. For each of these assays, Peripheral blood mononuclear cells (PBMC) are purified from single donor leukopacks (American Red Cross, Baltimore, MD) by centrifugation through a Histopaque gradient (Sigma). Monocytes are isolated from PBMC by counterflow centrifugal elutriation.

Monocyte Survival Assay. Human peripheral blood monocytes progressively lose viability when cultured in absence of serum or other stimuli. Their death results from

internally regulated process (apoptosis). Addition to the culture of activating factors, such as TNF-alpha dramatically improves cell survival and prevents DNA fragmentation. Propidium iodide (PI) staining is used to measure apoptosis as follows. Monocytes are cultured for 48 hours in polypropylene tubes in serum-free medium (positive control), in the presence of 100 ng/ml TNF-alpha (negative control), and in the presence of varying concentrations of the compound to be tested. Cells are suspended at a concentration of 2×10^6 /ml in PBS containing PI at a final concentration of 5 µg/ml, and then incubated at room temperature for 5 minutes before FACScan analysis. PI uptake has been demonstrated to correlate with DNA fragmentation in this experimental paradigm.

Effect on cytokine release. An important function of monocytes/macrophages is their regulatory activity on other cellular populations of the immune system through the release of cytokines after stimulation. An ELISA to measure cytokine release is performed as follows. Human monocytes are incubated at a density of 5×10^5 cells/ml with increasing concentrations of agonists or antagonists of the invention and under the same conditions, but in the absence of agonists or antagonists. For IL-12 production, the cells are primed overnight with IFN (100 U/ml) in presence of agonist or antagonist of the invention. LPS (10 ng/ml) is then added. Conditioned media are collected after 24h and kept frozen until use. Measurement of TNF-alpha, IL-10, MCP-1 and IL-8 is then performed using a commercially available ELISA kit (e. g, R & D Systems (Minneapolis, MN)) and applying the standard protocols provided with the kit.

Oxidative burst. Purified monocytes are plated in 96-w plate at 2×10^5 cell/well. Increasing concentrations of agonists or antagonists of the invention are added to the wells in a total volume of 0.2 ml culture medium (RPMI 1640 + 10% FCS, glutamine and antibiotics). After 3 days incubation, the plates are centrifuged and the medium is removed from the wells. To the macrophage monolayers, 0.2 ml per well of phenol red solution (140 mM NaCl, 10 mM potassium phosphate buffer pH 7.0, 5.5 mM dextrose, 0.56 mM phenol red and 19 U/ml of HRPO) is added, together with the stimulant (200 nM PMA). The plates are incubated at 37°C for 2 hours and the reaction is stopped by adding 20 µl 1N NaOH per well. The absorbance is read at 610 nm. To calculate the amount of H_2O_2 produced by the

macrophages, a standard curve of a H₂O₂ solution of known molarity is performed for each experiment.

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

Example 25: Biological Effects of Agonists or Antagonists of the Invention

Astrocyte and Neuronal Assays

Agonists or antagonists of the invention, expressed in *Escherichia coli* and purified as described above, can be tested for activity in promoting the survival, neurite outgrowth, or phenotypic differentiation of cortical neuronal cells and for inducing the proliferation of glial fibrillary acidic protein immunopositive cells, astrocytes. The selection of cortical cells for the bioassay is based on the prevalent expression of FGF-1 and FGF-2 in cortical structures and on the previously reported enhancement of cortical neuronal survival resulting from FGF-2 treatment. A thymidine incorporation assay, for example, can be used to elucidate an agonist or antagonist of the invention's activity on these cells.

Moreover, previous reports describing the biological effects of FGF-2 (basic FGF) on cortical or hippocampal neurons *in vitro* have demonstrated increases in both neuron survival and neurite outgrowth (Walicke et al., "Fibroblast growth factor promotes survival of dissociated hippocampal neurons and enhances neurite extension." *Proc. Natl. Acad. Sci. USA* 83:3012-3016. (1986), assay herein incorporated by reference in its entirety). However, reports from experiments done on PC-12 cells suggest that these two responses are not necessarily synonymous and may depend on not only which FGF is being tested but also on which receptor(s) are expressed on the target cells. Using the primary cortical neuronal culture paradigm, the ability of an agonist or antagonist of the invention to induce neurite outgrowth can be compared to the response achieved with FGF-2 using, for example, a thymidine incorporation assay.

Fibroblast and endothelial cell assays

Human lung fibroblasts are obtained from Clonetics (San Diego, CA) and maintained in growth media from Clonetics. Dermal microvascular endothelial cells are obtained from Cell Applications (San Diego, CA). For proliferation assays, the human lung fibroblasts and dermal microvascular endothelial cells can be cultured at 5,000 cells/well in a 96-well plate for one day in growth medium. The cells are then incubated for one day in 0.1% BSA basal medium. After replacing the medium with fresh 0.1% BSA medium, the cells are incubated with the test proteins for 3 days. Alamar Blue (Alamar Biosciences, Sacramento, CA) is added to each well to a final concentration of 10%. The cells are incubated for 4 hr. Cell viability is measured by reading in a CytoFluor fluorescence reader. For the PGE₂ assays, the human lung fibroblasts are cultured at 5,000 cells/well in a 96-well plate for one day. After a medium change to 0.1% BSA basal medium, the cells are incubated with FGF-2 or agonists or antagonists of the invention with or without IL-1 α for 24 hours. The supernatants are collected and assayed for PGE₂ by EIA kit (Cayman, Ann Arbor, MI). For the IL-6 assays, the human lung fibroblasts are cultured at 5,000 cells/well in a 96-well plate for one day. After a medium change to 0.1% BSA basal medium, the cells are incubated with FGF-2 or with or without agonists or antagonists of the invention IL-1 α for 24 hours. The supernatants are collected and assayed for IL-6 by ELISA kit (Endogen, Cambridge, MA).

Human lung fibroblasts are cultured with FGF-2 or agonists or antagonists of the invention for 3 days in basal medium before the addition of Alamar Blue to assess effects on growth of the fibroblasts. FGF-2 should show a stimulation at 10 - 2500 ng/ml which can be used to compare stimulation with agonists or antagonists of the invention.

Parkinson Models.

The loss of motor function in Parkinson's disease is attributed to a deficiency of striatal dopamine resulting from the degeneration of the nigrostriatal dopaminergic projection neurons. An animal model for Parkinson's that has been extensively characterized involves the systemic administration of 1-methyl-4 phenyl 1,2,3,6-tetrahydropyridine (MPTP). In the CNS, MPTP is taken-up by astrocytes and catabolized by monoamine oxidase B to 1-methyl-4-phenyl pyridine (MPP⁺) and released. Subsequently, MPP⁺ is actively accumulated in dopaminergic neurons by the high-affinity reuptake transporter for dopamine. MPP⁺ is then

concentrated in mitochondria by the electrochemical gradient and selectively inhibits nicotinamide adenine disphosphate: ubiquinone oxidoreductionase (complex I), thereby interfering with electron transport and eventually generating oxygen radicals.

It has been demonstrated in tissue culture paradigms that FGF-2 (basic FGF) has trophic activity towards nigral dopaminergic neurons (Ferrari et al., Dev. Biol. 1989). Recently, Dr. Unsicker's group has demonstrated that administering FGF-2 in gel foam implants in the striatum results in the near complete protection of nigral dopaminergic neurons from the toxicity associated with MPTP exposure (Otto and Unsicker, J. Neuroscience, 1990).

Based on the data with FGF-2, agonists or antagonists of the invention can be evaluated to determine whether it has an action similar to that of FGF-2 in enhancing dopaminergic neuronal survival *in vitro* and it can also be tested *in vivo* for protection of dopaminergic neurons in the striatum from the damage associated with MPTP treatment. The potential effect of an agonist or antagonist of the invention is first examined *in vitro* in a dopaminergic neuronal cell culture paradigm. The cultures are prepared by dissecting the midbrain floor plate from gestation day 14 Wistar rat embryos. The tissue is dissociated with trypsin and seeded at a density of 200,000 cells/cm² on polyorthinine-laminin coated glass coverslips. The cells are maintained in Dulbecco's Modified Eagle's medium and F12 medium containing hormonal supplements (N1). The cultures are fixed with paraformaldehyde after 8 days *in vitro* and are processed for tyrosine hydroxylase, a specific marker for dopaminergic neurons, immunohistochemical staining. Dissociated cell cultures are prepared from embryonic rats. The culture medium is changed every third day and the factors are also added at that time.

Since the dopaminergic neurons are isolated from animals at gestation day 14, a developmental time which is past the stage when the dopaminergic precursor cells are proliferating, an increase in the number of tyrosine hydroxylase immunopositive neurons would represent an increase in the number of dopaminergic neurons surviving *in vitro*. Therefore, if an agonist or antagonist of the invention acts to prolong the survival of dopaminergic neurons, it would suggest that the agonist or antagonist may be involved in Parkinson's Disease.

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

5

Example 26: The Effect of Agonists or Antagonists of the Invention on the Growth of Vascular Endothelial Cells

On day 1, human umbilical vein endothelial cells (HUVEC) are seeded at $2-5 \times 10^4$ cells/35 mm dish density in M199 medium containing 4% fetal bovine serum (FBS), 16 units/ml heparin, and 50 units/ml endothelial cell growth supplements (ECGS, Biotechnology, Inc.). On day 2, the medium is replaced with M199 containing 10% FBS, 8 units/ml heparin. An agonist or antagonist of the invention, and positive controls, such as VEGF and basic FGF (bFGF) are added, at varying concentrations. On days 4 and 6, the medium is replaced. On day 8, cell number is determined with a Coulter Counter.

15 An increase in the number of HUVEC cells indicates that the compound of the invention may proliferate vascular endothelial cells, while a decrease in the number of HUVEC cell indicates that the compound of the invention inhibits vascular endothelial cells.

The studies described in this example tested activity of a polypeptide of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides (e.g., gene therapy), agonists, and/or antagonists of the invention.

20

Example 27: Rat Corneal Wound Healing Model

This animal model shows the effect of an agonist or antagonist of the invention on neovascularization. The experimental protocol includes:

- 25 a) Making a 1-1.5 mm long incision from the center of cornea into the stromal layer.
- b) Inserting a spatula below the lip of the incision facing the outer corner of the eye.
- c) Making a pocket (its base is 1-1.5 mm from the edge of the eye).
- 30 d) Positioning a pellet, containing 50ng- 5ug of an agonist or antagonist of the invention, within the pocket.

e) Treatment with an agonist or antagonist of the invention can also be applied topically to the corneal wounds in a dosage range of 20mg - 500mg (daily treatment for five days).

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

Example 28: Diabetic Mouse and Glucocorticoid-Impaired Wound Healing Models

A. Diabetic db+/db+ Mouse Model.

To demonstrate that an agonist or antagonist of the invention accelerates the healing process, the genetically diabetic mouse model of wound healing is used. The full thickness wound healing model in the db+/db+ mouse is a well characterized, clinically relevant and reproducible model of impaired wound healing. Healing of the diabetic wound is dependent on formation of granulation tissue and re-epithelialization rather than contraction (Gartner, M.H. *et al.*, *J. Surg. Res.* 52:389 (1992); Greenhalgh, D.G. *et al.*, *Am. J. Pathol.* 136:1235 (1990)).

The diabetic animals have many of the characteristic features observed in Type II diabetes mellitus. Homozygous (db+/db+) mice are obese in comparison to their normal heterozygous (db+/+m) littermates. Mutant diabetic (db+/db+) mice have a single autosomal recessive mutation on chromosome 4 (db+) (Coleman *et al.* *Proc. Natl. Acad. Sci. USA* 77:283-293 (1982)). Animals show polyphagia, polydipsia and polyuria. Mutant diabetic mice (db+/db+) have elevated blood glucose, increased or normal insulin levels, and suppressed cell-mediated immunity (Mandel *et al.*, *J. Immunol.* 120:1375 (1978); Debray-Sachs, M. *et al.*, *Clin. Exp. Immunol.* 51(1):1-7 (1983); Leiter *et al.*, *Am. J. of Pathol.* 114:46-55 (1985)). Peripheral neuropathy, myocardial complications, and microvascular lesions, basement membrane thickening and glomerular filtration abnormalities have been described in these animals (Norido, F. *et al.*, *Exp. Neurol.* 83(2):221-232 (1984); Robertson *et al.*, *Diabetes* 29(1):60-67 (1980); Giacomelli *et al.*, *Lab Invest.* 40(4):460-473 (1979); Coleman, D.L., *Diabetes* 31 (Suppl):1-6 (1982)). These homozygous diabetic mice develop hyperglycemia that is resistant to insulin analogous to human type II diabetes (Mandel *et al.*, *J. Immunol.* 120:1375-1377 (1978)).

The characteristics observed in these animals suggests that healing in this model may be similar to the healing observed in human diabetes (Greenhalgh, *et al.*, *Am. J. of Pathol.* 136:1235-1246 (1990)).

Genetically diabetic female C57BL/KsJ (db+/db+) mice and their non-diabetic (db+/+m) heterozygous littermates are used in this study (Jackson Laboratories). The animals are purchased at 6 weeks of age and are 8 weeks old at the beginning of the study. Animals are individually housed and received food and water ad libitum. All manipulations are performed using aseptic techniques. The experiments are conducted according to the rules and guidelines of Human Genome Sciences, Inc. Institutional Animal Care and Use Committee and the Guidelines for the Care and Use of Laboratory Animals.

Wounding protocol is performed according to previously reported methods (Tsuboi, R. and Rifkin, D.B., *J. Exp. Med.* 172:245-251 (1990)). Briefly, on the day of wounding, animals are anesthetized with an intraperitoneal injection of Avertin (0.01 mg/mL), 2,2,2-tribromoethanol and 2-methyl-2-butanol dissolved in deionized water. The dorsal region of the animal is shaved and the skin washed with 70% ethanol solution and iodine. The surgical area is dried with sterile gauze prior to wounding. An 8 mm full-thickness wound is then created using a Keyes tissue punch. Immediately following wounding, the surrounding skin is gently stretched to eliminate wound expansion. The wounds are left open for the duration of the experiment. Application of the treatment is given topically for 5 consecutive days commencing on the day of wounding. Prior to treatment, wounds are gently cleansed with sterile saline and gauze sponges.

Wounds are visually examined and photographed at a fixed distance at the day of surgery and at two day intervals thereafter. Wound closure is determined by daily measurement on days 1-5 and on day 8. Wounds are measured horizontally and vertically using a calibrated Jameson caliper. Wounds are considered healed if granulation tissue is no longer visible and the wound is covered by a continuous epithelium.

An agonist or antagonist of the invention is administered using at a range different doses, from 4mg to 500mg per wound per day for 8 days in vehicle. Vehicle control groups received 50mL of vehicle solution.

Animals are euthanized on day 8 with an intraperitoneal injection of sodium pentobarbital (300mg/kg). The wounds and surrounding skin are then harvested for histology

and immunohistochemistry. Tissue specimens are placed in 10% neutral buffered formalin in tissue cassettes between biopsy sponges for further processing.

Three groups of 10 animals each (5 diabetic and 5 non-diabetic controls) are evaluated: 1) Vehicle placebo control, 2) untreated group, and 3) treated group.

5 Wound closure is analyzed by measuring the area in the vertical and horizontal axis and obtaining the total square area of the wound. Contraction is then estimated by establishing the differences between the initial wound area (day 0) and that of post treatment (day 8). The wound area on day 1 is 64mm², the corresponding size of the dermal punch. Calculations are made using the following formula:

10
$$[\text{Open area on day 8}] - [\text{Open area on day 1}] / [\text{Open area on day 1}]$$

Specimens are fixed in 10% buffered formalin and paraffin embedded blocks are sectioned perpendicular to the wound surface (5mm) and cut using a Reichert-Jung
15 microtome. Routine hematoxylin-eosin (H&E) staining is performed on cross-sections of bisected wounds. Histologic examination of the wounds are used to assess whether the healing process and the morphologic appearance of the repaired skin is altered by treatment with an agonist or antagonist of the invention. This assessment included verification of the presence of cell accumulation, inflammatory cells, capillaries, fibroblasts, re-epithelialization
20 and epidermal maturity (Greenhalgh, D.G. *et al.*, *Am. J. Pathol.* 136:1235 (1990)). A calibrated lens micrometer is used by a blinded observer.

Tissue sections are also stained immunohistochemically with a polyclonal rabbit anti-human keratin antibody using ABC Elite detection system. Human skin is used as a positive tissue control while non-immune IgG is used as a negative control. Keratinocyte growth is
25 determined by evaluating the extent of reepithelialization of the wound using a calibrated lens micrometer.

Proliferating cell nuclear antigen/cyclin (PCNA) in skin specimens is demonstrated by using anti-PCNA antibody (1:50) with an ABC Elite detection system. Human colon cancer served as a positive tissue control and human brain tissue is used as a negative tissue control.
30 Each specimen included a section with omission of the primary antibody and substitution with non-immune mouse IgG. Ranking of these sections is based on the extent of

proliferation on a scale of 0-8, the lower side of the scale reflecting slight proliferation to the higher side reflecting intense proliferation.

Experimental data are analyzed using an unpaired t test. A p value of < 0.05 is considered significant.

5

B. Steroid Impaired Rat Model

The inhibition of wound healing by steroids has been well documented in various *in vitro* and *in vivo* systems (Wahl, Glucocorticoids and Wound healing. In: Anti-Inflammatory Steroid Action: Basic and Clinical Aspects. 280-302 (1989); Wahl *et al.*, *J. Immunol.* 115: 476-481 (1975); Werb *et al.*, *J. Exp. Med.* 147:1684-1694 (1978)). Glucocorticoids retard wound healing by inhibiting angiogenesis, decreasing vascular permeability (Ebert *et al.*, *Ann. Intern. Med.* 37:701-705 (1952)), fibroblast proliferation, and collagen synthesis (Beck *et al.*, *Growth Factors.* 5: 295-304 (1991); Haynes *et al.*, *J. Clin. Invest.* 61: 703-797 (1978)) and producing a transient reduction of circulating monocytes (Haynes *et al.*, *J. Clin. Invest.* 61: 703-797 (1978); Wahl, "Glucocorticoids and wound healing", In: Antiinflammatory Steroid Action: Basic and Clinical Aspects, Academic Press, New York, pp. 280-302 (1989)). The systemic administration of steroids to impaired wound healing is a well established phenomenon in rats (Beck *et al.*, *Growth Factors.* 5: 295-304 (1991); Haynes *et al.*, *J. Clin. Invest.* 61: 703-797 (1978); Wahl, "Glucocorticoids and wound healing", In: Antiinflammatory Steroid Action: Basic and Clinical Aspects, Academic Press, New York, pp. 280-302 (1989); Pierce *et al.*, *Proc. Natl. Acad. Sci. USA* 86: 2229-2233 (1989)).

To demonstrate that an agonist or antagonist of the invention can accelerate the healing process, the effects of multiple topical applications of the agonist or antagonist on full thickness excisional skin wounds in rats in which healing has been impaired by the systemic administration of methylprednisolone is assessed.

Young adult male Sprague Dawley rats weighing 250-300 g (Charles River Laboratories) are used in this example. The animals are purchased at 8 weeks of age and are 9 weeks old at the beginning of the study. The healing response of rats is impaired by the systemic administration of methylprednisolone (17mg/kg/rat intramuscularly) at the time of wounding. Animals are individually housed and received food and water *ad libitum*. All manipulations are performed using aseptic techniques. This study is conducted according to

the rules and guidelines of Human Genome Sciences, Inc. Institutional Animal Care and Use Committee and the Guidelines for the Care and Use of Laboratory Animals.

The wounding protocol is followed according to section A, above. On the day of wounding, animals are anesthetized with an intramuscular injection of ketamine (50 mg/kg) and xylazine (5 mg/kg). The dorsal region of the animal is shaved and the skin washed with 70% ethanol and iodine solutions. The surgical area is dried with sterile gauze prior to wounding. An 8 mm full-thickness wound is created using a Keyes tissue punch. The wounds are left open for the duration of the experiment. Applications of the testing materials are given topically once a day for 7 consecutive days commencing on the day of wounding and subsequent to methylprednisolone administration. Prior to treatment, wounds are gently cleansed with sterile saline and gauze sponges.

Wounds are visually examined and photographed at a fixed distance at the day of wounding and at the end of treatment. Wound closure is determined by daily measurement on days 1-5 and on day 8. Wounds are measured horizontally and vertically using a calibrated Jameson caliper. Wounds are considered healed if granulation tissue is no longer visible and the wound is covered by a continuous epithelium.

The agonist or antagonist of the invention is administered using at a range different doses, from 4mg to 500mg per wound per day for 8 days in vehicle. Vehicle control groups received 50mL of vehicle solution.

Animals are euthanized on day 8 with an intraperitoneal injection of sodium pentobarbital (300mg/kg). The wounds and surrounding skin are then harvested for histology. Tissue specimens are placed in 10% neutral buffered formalin in tissue cassettes between biopsy sponges for further processing.

Four groups of 10 animals each (5 with methylprednisolone and 5 without glucocorticoid) are evaluated: 1) Untreated group 2) Vehicle placebo control 3) treated groups.

Wound closure is analyzed by measuring the area in the vertical and horizontal axis and obtaining the total area of the wound. Closure is then estimated by establishing the differences between the initial wound area (day 0) and that of post treatment (day 8). The wound area on day 1 is 64mm^2 , the corresponding size of the dermal punch. Calculations are made using the following formula:

[Open area on day 8] - [Open area on day 1] / [Open area on day 1]

Specimens are fixed in 10% buffered formalin and paraffin embedded blocks are sectioned perpendicular to the wound surface (5mm) and cut using an Olympus microtome.

5 Routine hematoxylin-eosin (H&E) staining is performed on cross-sections of bisected wounds. Histologic examination of the wounds allows assessment of whether the healing process and the morphologic appearance of the repaired skin is improved by treatment with an agonist or antagonist of the invention. A calibrated lens micrometer is used by a blinded observer to determine the distance of the wound gap.

10 Experimental data are analyzed using an unpaired t test. A p value of < 0.05 is considered significant.

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

15

Example 29: Lymphadema Animal Model

The purpose of this experimental approach is to create an appropriate and consistent lymphedema model for testing the therapeutic effects of an agonist or antagonist of the invention in lymphangiogenesis and re-establishment of the lymphatic circulatory system in
20 the rat hind limb. Effectiveness is measured by swelling volume of the affected limb, quantification of the amount of lymphatic vasculature, total blood plasma protein, and histopathology. Acute lymphedema is observed for 7-10 days. Perhaps more importantly, the chronic progress of the edema is followed for up to 3-4 weeks.

Prior to beginning surgery, blood sample is drawn for protein concentration analysis.
25 Male rats weighing approximately ~350g are dosed with Pentobarbital. Subsequently, the right legs are shaved from knee to hip. The shaved area is swabbed with gauze soaked in 70% EtOH. Blood is drawn for serum total protein testing. Circumference and volumetric measurements are made prior to injecting dye into paws after marking 2 measurement levels (0.5 cm above heel, at mid-pt of dorsal paw). The intradermal dorsum of both right and left
30 paws are injected with 0.05 ml of 1% Evan's Blue. Circumference and volumetric measurements are then made following injection of dye into paws.

Using the knee joint as a landmark, a mid-leg inguinal incision is made circumferentially allowing the femoral vessels to be located. Forceps and hemostats are used to dissect and separate the skin flaps. After locating the femoral vessels, the lymphatic vessel that runs along side and underneath the vessel(s) is located. The main lymphatic vessels in this area are then electrically coagulated or suture ligated.

Using a microscope, muscles in back of the leg (near the semitendinosus and adductors) are bluntly dissected. The popliteal lymph node is then located. The 2 proximal and 2 distal lymphatic vessels and distal blood supply of the popliteal node are then and ligated by suturing. The popliteal lymph node, and any accompanying adipose tissue, is then removed by cutting connective tissues.

Care is taken to control any mild bleeding resulting from this procedure. After lymphatics are occluded, the skin flaps are sealed by using liquid skin (Vetbond) (AJ Buck). The separated skin edges are sealed to the underlying muscle tissue while leaving a gap of ~0.5 cm around the leg. Skin also may be anchored by suturing to underlying muscle when necessary.

To avoid infection, animals are housed individually with mesh (no bedding). Recovering animals are checked daily through the optimal edematous peak, which typically occurred by day 5-7. The plateau edematous peak are then observed. To evaluate the intensity of the lymphedema, the circumference and volumes of 2 designated places on each paw before operation and daily for 7 days are measured. The effect plasma proteins on lymphedema is determined and whether protein analysis is a useful testing perimeter is also investigated. The weights of both control and edematous limbs are evaluated at 2 places. Analysis is performed in a blind manner.

Circumference Measurements: Under brief gas anesthetic to prevent limb movement, a cloth tape is used to measure limb circumference. Measurements are done at the ankle bone and dorsal paw by 2 different people then those 2 readings are averaged. Readings are taken from both control and edematous limbs.

Volumetric Measurements: On the day of surgery, animals are anesthetized with Pentobarbital and are tested prior to surgery. For daily volumetrics animals are under brief halothane anesthetic (rapid immobilization and quick recovery), both legs are shaved and equally marked using waterproof marker on legs. Legs are first dipped in water, then dipped

into instrument to each marked level then measured by Buxco edema software(Chen/Victor). Data is recorded by one person, while the other is dipping the limb to marked area.

Blood-plasma protein measurements: Blood is drawn, spun, and serum separated prior to surgery and then at conclusion for total protein and Ca²⁺ comparison.

5 Limb Weight Comparison: After drawing blood, the animal is prepared for tissue collection. The limbs are amputated using a quillitine, then both experimental and control legs are cut at the ligature and weighed. A second weighing is done as the tibio-cacaneal joint is disarticulated and the foot is weighed.

10 Histological Preparations: The transverse muscle located behind the knee (popliteal) area is dissected and arranged in a metal mold, filled with freezeGel, dipped into cold methylbutane, placed into labeled sample bags at - 80EC until sectioning. Upon sectioning, the muscle is observed under fluorescent microscopy for lymphatics..

15 The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

Example 30: Suppression of TNF alpha-induced adhesion molecule expression by a Agonist or Antagonist of the Invention

20 The recruitment of lymphocytes to areas of inflammation and angiogenesis involves specific receptor-ligand interactions between cell surface adhesion molecules (CAMs) on lymphocytes and the vascular endothelium. The adhesion process, in both normal and pathological settings, follows a multi-step cascade that involves intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion molecule-1 (VCAM-1), and endothelial leukocyte adhesion molecule-1 (E-selectin) expression on endothelial cells (EC). The
25 expression of these molecules and others on the vascular endothelium determines the efficiency with which leukocytes may adhere to the local vasculature and extravasate into the local tissue during the development of an inflammatory response. The local concentration of cytokines and growth factor participate in the modulation of the expression of these CAMs.

30 Tumor necrosis factor alpha (TNF-a), a potent proinflammatory cytokine, is a stimulator of all three CAMs on endothelial cells and may be involved in a wide variety of inflammatory responses, often resulting in a pathological outcome.

The potential of an agonist or antagonist of the invention to mediate a suppression of TNF- α induced CAM expression can be examined. A modified ELISA assay which uses ECs as a solid phase absorbent is employed to measure the amount of CAM expression on TNF- α treated ECs when co-stimulated with a member of the FGF family of proteins.

5 To perform the experiment, human umbilical vein endothelial cell (HUVEC) cultures are obtained from pooled cord harvests and maintained in growth medium (EGM-2; Clonetics, San Diego, CA) supplemented with 10% FCS and 1% penicillin/streptomycin in a 37 degree C humidified incubator containing 5% CO₂. HUVECs are seeded in 96-well plates at concentrations of 1×10^4 cells/well in EGM medium at 37 degree C for 18-24 hrs or
10 until confluent. The monolayers are subsequently washed 3 times with a serum-free solution of RPMI-1640 supplemented with 100 U/ml penicillin and 100 mg/ml streptomycin, and treated with a given cytokine and/or growth factor(s) for 24 h at 37 degree C. Following incubation, the cells are then evaluated for CAM expression.

Human Umbilical Vein Endothelial cells (HUVECs) are grown in a standard 96 well
15 plate to confluence. Growth medium is removed from the cells and replaced with 90 μ l of 199 Medium (10% FBS). Samples for testing and positive or negative controls are added to the plate in triplicate (in 10 μ l volumes). Plates are incubated at 37 degree C for either 5 h (selectin and integrin expression) or 24 h (integrin expression only). Plates are aspirated to remove medium and 100 μ l of 0.1% paraformaldehyde-PBS(with Ca⁺⁺ and Mg⁺⁺) is added
20 to each well. Plates are held at 4°C for 30 min.

Fixative is then removed from the wells and wells are washed 1X with PBS(+Ca,Mg)+0.5% BSA and drained. Do not allow the wells to dry. Add 10 μ l of diluted primary antibody to the test and control wells. Anti-ICAM-1-Biotin, Anti-VCAM-1-Biotin and Anti-E-selectin-Biotin are used at a concentration of 10 μ g/ml (1:10 dilution of 0.1
25 mg/ml stock antibody). Cells are incubated at 37°C for 30 min. in a humidified environment. Wells are washed X3 with PBS(+Ca,Mg)+0.5% BSA.

Then add 20 μ l of diluted ExtrAvidin-Alkaline Phosphatase (1:5,000 dilution) to each well and incubated at 37°C for 30 min. Wells are washed X3 with PBS(+Ca,Mg)+0.5% BSA. 1 tablet of p-Nitrophenol Phosphate pNPP is dissolved in 5 ml of glycine buffer (pH
30 10.4). 100 μ l of pNPP substrate in glycine buffer is added to each test well. Standard wells in triplicate are prepared from the working dilution of the ExtrAvidin-Alkaline Phosphatase in glycine buffer: $1:5,000 (10^0) > 10^{-0.5} > 10^{-1} > 10^{-1.5}$. 5 μ l of each dilution is added to triplicate

wells and the resulting AP content in each well is 5.50 ng, 1.74 ng, 0.55 ng, 0.18 ng. 100 μ l of pNPNP reagent must then be added to each of the standard wells. The plate must be incubated at 37°C for 4h. A volume of 50 μ l of 3M NaOH is added to all wells. The results are quantified on a plate reader at 405 nm. The background subtraction option is used on blank wells filled with glycine buffer only. The template is set up to indicate the concentration of AP-conjugate in each standard well [5.50 ng; 1.74 ng; 0.55 ng; 0.18 ng]. Results are indicated as amount of bound AP-conjugate in each sample.

The studies described in this example tested activity of agonists or antagonists of the invention. However, one skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides or polypeptides of the invention (e.g., gene therapy).

Example 31: Production Of Polypeptide of the Invention For High-Throughput Screening Assays.

The following protocol produces a supernatant containing polypeptide of the present invention to be tested. This supernatant can then be used in the Screening Assays described in Examples 33-42.

First, dilute Poly-D-Lysine (644 587 Boehringer-Mannheim) stock solution (1mg/ml in PBS) 1:20 in PBS (w/o calcium or magnesium 17-516F Biowhittaker) for a working solution of 50ug/ml. Add 200 μ l of this solution to each well (24 well plates) and incubate at RT for 20 minutes. Be sure to distribute the solution over each well (note: a 12-channel pipetter may be used with tips on every other channel). Aspirate off the Poly-D-Lysine solution and rinse with 1ml PBS (Phosphate Buffered Saline). The PBS should remain in the well until just prior to plating the cells and plates may be poly-lysine coated in advance for up to two weeks.

Plate 293T cells (do not carry cells past P+20) at 2×10^5 cells/well in .5ml DMEM(Dulbecco's Modified Eagle Medium)(with 4.5 G/L glucose and L-glutamine (12-604F Biowhittaker))/10% heat inactivated FBS(14-503F Biowhittaker)/1x Penstrep(17-602E Biowhittaker). Let the cells grow overnight.

The next day, mix together in a sterile solution basin: 300 μ l Lipofectamine (18324-012 Gibco/BRL) and 5ml Optimem I (31985070 Gibco/BRL)/96-well plate. With a small volume multi-channel pipetter, aliquot approximately 2ug of an expression vector containing a polynucleotide insert, produced by the methods described in Examples 8-10, into an

appropriately labeled 96-well round bottom plate. With a multi-channel pipetter, add 50ul of the Lipofectamine/Optimem I mixture to each well. Pipette up and down gently to mix. Incubate at RT 15-45 minutes. After about 20 minutes, use a multi-channel pipetter to add 150ul Optimem I to each well. As a control, one plate of vector DNA lacking an insert
5 should be transfected with each set of transfections.

Preferably, the transfection should be performed by tag-teaming the following tasks. By tag-teaming, hands on time is cut in half, and the cells do not spend too much time on PBS. First, person A aspirates off the media from four 24-well plates of cells, and then person B rinses each well with .5-1ml PBS. Person A then aspirates off PBS rinse, and
10 person B, using a 12-channel pipetter with tips on every other channel, adds the 200ul of DNA/Lipofectamine/Optimem I complex to the odd wells first, then to the even wells, to each row on the 24-well plates. Incubate at 37 degree C for 6 hours.

While cells are incubating, prepare appropriate media, either 1%BSA in DMEM with 1x penstrep, or HGS CHO-5 media (116.6 mg/L of CaCl_2 (anhyd); 0.00130 mg/L $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$; 0.050 mg/L of $\text{Fe}(\text{NO}_3)_3 \cdot 9\text{H}_2\text{O}$; 0.417 mg/L of $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$; 311.80 mg/L of KCl; 28.64 mg/L of MgCl_2 ; 48.84 mg/L of MgSO_4 ; 6995.50 mg/L of NaCl; 2400.0 mg/L of NaHCO_3 ; 62.50 mg/L of $\text{NaH}_2\text{PO}_4 \cdot \text{H}_2\text{O}$; 71.02 mg/L of Na_2HPO_4 ; .4320 mg/L of $\text{ZnSO}_4 \cdot 7\text{H}_2\text{O}$; .002 mg/L of Arachidonic Acid ; 1.022 mg/L of Cholesterol; .070 mg/L of DL-alpha-Tocopherol-Acetate; 0.0520 mg/L of Linoleic Acid; 0.010 mg/L of Linolenic Acid; 0.010
20 mg/L of Myristic Acid; 0.010 mg/L of Oleic Acid; 0.010 mg/L of Palmitic Acid; 0.010 mg/L of Palmitic Acid; 100 mg/L of Pluronic F-68; 0.010 mg/L of Stearic Acid; 2.20 mg/L of Tween 80; 4551 mg/L of D-Glucose; 130.85 mg/ml of L- Alanine; 147.50 mg/ml of L- Arginine-HCL; 7.50 mg/ml of L-Asparagine- H_2O ; 6.65 mg/ml of L-Aspartic Acid; 29.56 mg/ml of L-Cystine-2HCL- H_2O ; 31.29 mg/ml of L-Cystine-2HCL; 7.35 mg/ml of L-
25 Glutamic Acid; 365.0 mg/ml of L-Glutamine; 18.75 mg/ml of Glycine; 52.48 mg/ml of L-Histidine-HCL- H_2O ; 106.97 mg/ml of L-Isoleucine; 111.45 mg/ml of L-Leucine; 163.75 mg/ml of L-Lysine HCL; 32.34 mg/ml of L-Methionine; 68.48 mg/ml of L-Phenylalanine; 40.0 mg/ml of L-Proline; 26.25 mg/ml of L-Serine; 101.05 mg/ml of L-Threonine; 19.22 mg/ml of L-Tryptophan; 91.79 mg/ml of L-Tyrosine-2Na-2 H_2O ; and 99.65 mg/ml of L-
30 Valine; 0.0035 mg/L of Biotin; 3.24 mg/L of D-Ca Pantothenate; 11.78 mg/L of Choline

Chloride; 4.65 mg/L of Folic Acid; 15.60 mg/L of i-Inositol; 3.02 mg/L of Niacinamide; 3.00 mg/L of Pyridoxal HCL; 0.031 mg/L of Pyridoxine HCL; 0.319 mg/L of Riboflavin; 3.17 mg/L of Thiamine HCL; 0.365 mg/L of Thymidine; 0.680 mg/L of Vitamin B₁₂; 25 mM of HEPES Buffer; 2.39 mg/L of Na Hypoxanthine; 0.105 mg/L of Lipoic Acid; 0.081 mg/L of Sodium Putrescine-2HCL; 55.0 mg/L of Sodium Pyruvate; 0.0067 mg/L of Sodium Selenite; 20uM of Ethanolamine; 0.122 mg/L of Ferric Citrate; 41.70 mg/L of Methyl-B-Cyclodextrin complexed with Linoleic Acid; 33.33 mg/L of Methyl-B-Cyclodextrin complexed with Oleic Acid; 10 mg/L of Methyl-B-Cyclodextrin complexed with Retinal Acetate. Adjust osmolarity to 327 mOsm) with 2mm glutamine and 1x penstrep. (BSA (81-068-3 Bayer) 100gm dissolved in 1L DMEM for a 10% BSA stock solution). Filter the media and collect 50 ul for endotoxin assay in 15ml polystyrene conical.

The transfection reaction is terminated, preferably by tag-teaming, at the end of the incubation period. Person A aspirates off the transfection media, while person B adds 1.5ml appropriate media to each well. Incubate at 37 degree C for 45 or 72 hours depending on the media used: 1%BSA for 45 hours or CHO-5 for 72 hours.

On day four, using a 300ul multichannel pipetter, aliquot 600ul in one 1ml deep well plate and the remaining supernatant into a 2ml deep well. The supernatants from each well can then be used in the assays described in Examples 33-40.

It is specifically understood that when activity is obtained in any of the assays described below using a supernatant, the activity originates from either the polypeptide of the present invention directly (e.g., as a secreted protein) or by polypeptide of the present invention inducing expression of other proteins, which are then secreted into the supernatant. Thus, the invention further provides a method of identifying the protein in the supernatant characterized by an activity in a particular assay.

Example 32: Construction of GAS Reporter Construct

One signal transduction pathway involved in the differentiation and proliferation of cells is called the Jaks-STATs pathway. Activated proteins in the Jaks-STATs pathway bind to gamma activation site "GAS" elements or interferon-sensitive responsive element ("ISRE"), located in the promoter of many genes. The binding of a protein to these elements alter the expression of the associated gene.

GAS and ISRE elements are recognized by a class of transcription factors called Signal Transducers and Activators of Transcription, or "STATs." There are six members of the STATs family. Stat1 and Stat3 are present in many cell types, as is Stat2 (as response to IFN-alpha is widespread). Stat4 is more restricted and is not in many cell types though it has
5 been found in T helper class I, cells after treatment with IL-12. Stat5 was originally called mammary growth factor, but has been found at higher concentrations in other cells including myeloid cells. It can be activated in tissue culture cells by many cytokines.

The STATs are activated to translocate from the cytoplasm to the nucleus upon tyrosine phosphorylation by a set of kinases known as the Janus Kinase ("Jaks") family. Jaks
10 represent a distinct family of soluble tyrosine kinases and include Tyk2, Jak1, Jak2, and Jak3. These kinases display significant sequence similarity and are generally catalytically inactive in resting cells.

The Jaks are activated by a wide range of receptors summarized in the Table below. (Adapted from review by Schidler and Darnell, Ann. Rev. Biochem. 64:621-51 (1995).) A
15 cytokine receptor family, capable of activating Jaks, is divided into two groups: (a) Class 1 includes receptors for IL-2, IL-3, IL-4, IL-6, IL-7, IL-9, IL-11, IL-12, IL-15, Epo, PRL, GH, G-CSF, GM-CSF, LIF, CNTF, and thrombopoietin; and (b) Class 2 includes IFN-a, IFN-g, and IL-10. The Class 1 receptors share a conserved cysteine motif (a set of four conserved cysteines and one tryptophan) and a WSXWS motif (a membrane proximal region encoding
20 Trp-Ser-Xxx-Trp-Ser (SEQ ID NO:2)).

Thus, on binding of a ligand to a receptor, Jaks are activated, which in turn activate STATs, which then translocate and bind to GAS elements. This entire process is encompassed in the Jaks-STATs signal transduction pathway.

Therefore, activation of the Jaks-STATs pathway, reflected by the binding of the GAS
25 or the ISRE element, can be used to indicate proteins involved in the proliferation and differentiation of cells. For example, growth factors and cytokines are known to activate the Jaks-STATs pathway. (See Table below.) Thus, by using GAS elements linked to reporter molecules, activators of the Jaks-STATs pathway can be identified.

			<u>JAKs</u>				<u>STATS GAS(elements) or ISRE</u>
	<u>Ligand</u>	<u>tyk2</u>	<u>Jak1</u>	<u>Jak2</u>	<u>Jak3</u>		
	<u>IFN family</u>						
5	IFN-a/B	+	+	-	-	1,2,3	ISRE
	IFN-g		+	+	-	1	GAS (IRF1>Lys6>IFP)
	IL-10	+	?	?	-	1,3	
	<u>gp130 family</u>						
10	IL-6 (Pleiotrohic)	+	+	+	?	1,3	GAS (IRF1>Lys6>IFP)
	IL-11(Pleiotrohic)	?	+	?	?	1,3	
	OnM(Pleiotrohic)	?	+	+	?	1,3	
	LIF(Pleiotrohic)	?	+	+	?	1,3	
	CNTF(Pleiotrohic)	-/+	+	+	?	1,3	
15	G-CSF(Pleiotrohic)	?	+	?	?	1,3	
	IL-12(Pleiotrohic)	+	-	+	+	1,3	
	<u>g-C family</u>						
	IL-2 (lymphocytes)	-	+	-	+	1,3,5	GAS
20	IL-4 (lymph/myeloid)	-	+	-	+	6	GAS(IRF1=IFP>Ly6)(IgH)
	IL-7 (lymphocytes)	-	+	-	+	5	GAS
	IL-9 (lymphocytes)	-	+	-	+	5	GAS
	IL-13 (lymphocyte)	-	+	?	?	6	GAS
	IL-15	?	+	?	+	5	GAS
25	<u>gp140 family</u>						
	IL-3 (myeloid)	-	-	+	-	5	GAS (IRF1>IFP>>Ly6)
	IL-5 (myeloid)	-	-	+	-	5	GAS
	GM-CSF (myeloid)	-	-	+	-	5	GAS
30	<u>Growth hormone family</u>						
	GH	?	-	+	-	5	
	PRL	?	+/-	+	-	1,3,5	
	EPO	?	-	+	-	5	GAS(B-
35	CAS>IRF1=IFP>>Ly6)						
	<u>Receptor Tyrosine Kinases</u>						
	EGF	?	+	+	-	1,3	GAS (IRF1)
	PDGF	?	+	+	-	1,3	
40	CSF-1	?	+	+	-	1,3	GAS (not IRF1)

To construct a synthetic GAS containing promoter element, which is used in the Biological Assays described in Examples 33-34, a PCR based strategy is employed to generate a GAS-SV40 promoter sequence. The 5' primer contains four tandem copies of the GAS binding site found in the IRF1 promoter and previously demonstrated to bind STATs upon induction with a range of cytokines (Rothman et al., Immunity 1:457-468 (1994).), although other GAS or ISRE elements can be used instead. The 5' primer also contains 18bp of sequence complementary to the SV40 early promoter sequence and is flanked with an XhoI site. The sequence of the 5' primer is:

5':GCGCCTCGAGATTTCCCCGAAATCTAGATTTCCCCGAAATGATTTCCCCGAAAT
GATTTCCCCGAAATATCTGCCATCTCAATTAG:3' (SEQ ID NO:3)

The downstream primer is complementary to the SV40 promoter and is flanked with a Hind III site: 5':GCGGCAAGCTTTTTGCAAAGCCTAGGC:3' (SEQ ID NO:4)

PCR amplification is performed using the SV40 promoter template present in the B-gal:promoter plasmid obtained from Clontech. The resulting PCR fragment is digested with XhoI/Hind III and subcloned into BLSK2-. (Stratagene.) Sequencing with forward and reverse primers confirms that the insert contains the following sequence:

5':CTCGAGATTTCCCCGAAATCTAGATTTCCCCGAAATGATTTCCCCGAAATGATT
TCCCCGAAATATCTGCCATCTCAATTAGTCAGCAACCATAGTCCCGCCCCTAACT
CCGCCCATCCCGCCCCTAACTCCGCCCAGTTCCGCCCATTCTCCGCCCCATGGCTG
ACTAATTTTTTTTATTTATGCAGAGGCCGAGGCCGCCTCGGCCTCTGAGCTATTCC
AGAAGTAGTGAGGAGGCTTTTTTGGAGGCCCTAGGCTTTTGCAAAAAGCTT:3'
(SEQ ID NO:5)

With this GAS promoter element linked to the SV40 promoter, a GAS:SEAP2 reporter construct is next engineered. Here, the reporter molecule is a secreted alkaline phosphatase, or "SEAP." Clearly, however, any reporter molecule can be instead of SEAP, in this or in any of the other Examples. Well known reporter molecules that can be used instead of SEAP include chloramphenicol acetyltransferase (CAT), luciferase, alkaline phosphatase, B-galactosidase, green fluorescent protein (GFP), or any protein detectable by an antibody.

The above sequence confirmed synthetic GAS-SV40 promoter element is subcloned into the pSEAP-Promoter vector obtained from Clontech using HindIII and XhoI, effectively

replacing the SV40 promoter with the amplified GAS:SV40 promoter element, to create the GAS-SEAP vector. However, this vector does not contain a neomycin resistance gene, and therefore, is not preferred for mammalian expression systems.

Thus, in order to generate mammalian stable cell lines expressing the GAS-SEAP reporter, the GAS-SEAP cassette is removed from the GAS-SEAP vector using Sall and NotI, and inserted into a backbone vector containing the neomycin resistance gene, such as pGFP-1 (Clontech), using these restriction sites in the multiple cloning site, to create the GAS-SEAP/Neo vector. Once this vector is transfected into mammalian cells, this vector can then be used as a reporter molecule for GAS binding as described in Examples 33-34.

Other constructs can be made using the above description and replacing GAS with a different promoter sequence. For example, construction of reporter molecules containing NFK-B and EGR promoter sequences are described in Examples 35 and 36. However, many other promoters can be substituted using the protocols described in these Examples. For instance, SRE, IL-2, NFAT, or Osteocalcin promoters can be substituted, alone or in combination (e.g., GAS/NF-KB/EGR, GAS/NF-KB, IL-2/NFAT, or NF-KB/GAS). Similarly, other cell lines can be used to test reporter construct activity, such as HELA (epithelial), HUVEC (endothelial), Reh (B-cell), Saos-2 (osteoblast), HUVAC (aortic), or Cardiomyocyte.

Example 33: High-Throughput Screening Assay for T-cell Activity.

The following protocol is used to assess T-cell activity by identifying factors, and determining whether supernate containing a polypeptide of the invention proliferates and/or differentiates T-cells. T-cell activity is assessed using the GAS/SEAP/Neo construct produced in Example 32. Thus, factors that increase SEAP activity indicate the ability to activate the Jaks-STATS signal transduction pathway. The T-cell used in this assay is Jurkat T-cells (ATCC Accession No. TIB-152), although Molt-3 cells (ATCC Accession No. CRL-1552) and Molt-4 cells (ATCC Accession No. CRL-1582) cells can also be used.

Jurkat T-cells are lymphoblastic CD4⁺ Th1 helper cells. In order to generate stable cell lines, approximately 2 million Jurkat cells are transfected with the GAS-SEAP/neo vector using DMRIE-C (Life Technologies)(transfection procedure described below). The transfected cells are seeded to a density of approximately 20,000 cells per well and transfectants resistant to 1 mg/ml gentamicin selected. Resistant colonies are expanded and then

tested for their response to increasing concentrations of interferon gamma. The dose response of a selected clone is demonstrated.

Specifically, the following protocol will yield sufficient cells for 75 wells containing 200 ul of cells. Thus, it is either scaled up, or performed in multiple to generate sufficient
5 cells for multiple 96 well plates. Jurkat cells are maintained in RPMI + 10% serum with 1%Pen-Strep. Combine 2.5 mls of OPTI-MEM (Life Technologies) with 10 ug of plasmid DNA in a T25 flask. Add 2.5 ml OPTI-MEM containing 50 ul of DMRIE-C and incubate at room temperature for 15-45 mins.

During the incubation period, count cell concentration, spin down the required
10 number of cells (10^7 per transfection), and resuspend in OPTI-MEM to a final concentration of 10^7 cells/ml. Then add 1ml of 1×10^7 cells in OPTI-MEM to T25 flask and incubate at 37 degree C for 6 hrs. After the incubation, add 10 ml of RPMI + 15% serum.

The Jurkat:GAS-SEAP stable reporter lines are maintained in RPMI + 10% serum, 1 mg/ml Genticin, and 1% Pen-Strep. These cells are treated with supernatants containing
15 polypeptide of the present invention or polypeptide of the present invention induced polypeptides as produced by the protocol described in Example 31.

On the day of treatment with the supernatant, the cells should be washed and resuspended in fresh RPMI + 10% serum to a density of 500,000 cells per ml. The exact number of cells required will depend on the number of supernatants being screened. For one
20 96 well plate, approximately 10 million cells (for 10 plates, 100 million cells) are required.

Transfer the cells to a triangular reservoir boat, in order to dispense the cells into a 96 well dish, using a 12 channel pipette. Using a 12 channel pipette, transfer 200 ul of cells into each well (therefore adding 100, 000 cells per well).

After all the plates have been seeded, 50 ul of the supernatants are transferred directly
25 from the 96 well plate containing the supernatants into each well using a 12 channel pipette. In addition, a dose of exogenous interferon gamma (0.1, 1.0, 10 ng) is added to wells H9, H10, and H11 to serve as additional positive controls for the assay.

The 96 well dishes containing Jurkat cells treated with supernatants are placed in an incubator for 48 hrs (note: this time is variable between 48-72 hrs). 35 ul samples from each
30 well are then transferred to an opaque 96 well plate using a 12 channel pipette. The opaque plates should be covered (using sellophane covers) and stored at -20 degree C until SEAP assays are performed according to Example 37. The plates containing the remaining treated

cells are placed at 4 degree C and serve as a source of material for repeating the assay on a specific well if desired.

As a positive control, 100 Unit/ml interferon gamma can be used which is known to activate Jurkat T cells. Over 30 fold induction is typically observed in the positive control
5 wells.

The above protocol may be used in the generation of both transient, as well as, stable transfected cells, which would be apparent to those of skill in the art.

Example 34: High-Throughput Screening Assay Identifying Myeloid Activity

The following protocol is used to assess myeloid activity of polypeptide of the present invention by determining whether polypeptide of the present invention proliferates and/or differentiates myeloid cells. Myeloid cell activity is assessed using the GAS/SEAP/Neo construct produced in Example 32. Thus, factors that increase SEAP activity indicate the ability to activate the Jaks-STATS signal transduction pathway. The myeloid cell used in this assay is U937, a pre-monocyte cell line, although TF-1, HL60, or KG1 can be used.

To transiently transfect U937 cells with the GAS/SEAP/Neo construct produced in Example 32, a DEAE-Dextran method (Kharbanda et. al., 1994, Cell Growth & Differentiation, 5:259-265) is used. First, harvest 2×10^7 U937 cells and wash with PBS. The U937 cells are usually grown in RPMI 1640 medium containing 10% heat-inactivated fetal bovine serum (FBS) supplemented with 100 units/ml penicillin and 100 mg/ml streptomycin.

Next, suspend the cells in 1 ml of 20 mM Tris-HCl (pH 7.4) buffer containing 0.5 mg/ml DEAE-Dextran, 8 ug GAS-SEAP2-plasmid DNA, 140 mM NaCl, 5 mM KCl, 375 uM $\text{Na}_2\text{HPO}_4 \cdot 7\text{H}_2\text{O}$, 1 mM MgCl_2 , and 675 uM CaCl_2 . Incubate at 37 degrees C for 45 min.

Wash the cells with RPMI 1640 medium containing 10% FBS and then resuspend in 10 ml complete medium and incubate at 37 degree C for 36 hr.

The GAS-SEAP/U937 stable cells are obtained by growing the cells in 400 ug/ml G418. The G418-free medium is used for routine growth but every one to two months, the cells should be re-grown in 400 ug/ml G418 for couple of passages.

These cells are tested by harvesting 1×10^8 cells (this is enough for ten 96-well plates assay) and wash with PBS. Suspend the cells in 200 ml above described growth medium, with a final density of 5×10^5 cells/ml. Plate 200 ul cells per well in the 96-well plate (or 1×10^5 cells/well).

Add 50 ul of the supernatant prepared by the protocol described in Example 31. Incubate at 37 degree C for 48 to 72 hr. As a positive control, 100 Unit/ml interferon gamma can be used which is known to activate U937 cells. Over 30 fold induction is typically observed in the positive control wells. SEAP assay the supernatant according to the protocol described in Example 37.

Example 35: High-Throughput Screening Assay Identifying Neuronal Activity.

When cells undergo differentiation and proliferation, a group of genes are activated through many different signal transduction pathways. One of these genes, EGR1 (early growth response gene 1), is induced in various tissues and cell types upon activation. The promoter of EGR1 is responsible for such induction. Using the EGR1 promoter linked to reporter molecules, activation of cells can be assessed by polypeptide of the present invention.

Particularly, the following protocol is used to assess neuronal activity in PC12 cell lines. PC12 cells (rat phenochromocytoma cells) are known to proliferate and/or differentiate by activation with a number of mitogens, such as TPA (tetradecanoyl phorbol acetate), NGF (nerve growth factor), and EGF (epidermal growth factor). The EGR1 gene expression is activated during this treatment. Thus, by stably transfecting PC12 cells with a construct containing an EGR promoter linked to SEAP reporter, activation of PC12 cells by polypeptide of the present invention can be assessed.

The EGR/SEAP reporter construct can be assembled by the following protocol. The EGR-1 promoter sequence (-633 to +1)(Sakamoto K et al., Oncogene 6:867-871 (1991)) can be PCR amplified from human genomic DNA using the following primers:

5' GCGCTCGAGGGATGACAGCGATAGAACCCCGG -3' (SEQ ID NO:6)

5' GCGAAGCTTCGCGACTCCCCGGATCCGCCTC-3' (SEQ ID NO:7)

Using the GAS:SEAP/Neo vector produced in Example 32, EGR1 amplified product can then be inserted into this vector. Linearize the GAS:SEAP/Neo vector using restriction enzymes XhoI/HindIII, removing the GAS/SV40 stuffer. Restrict the EGR1 amplified product with these same enzymes. Ligate the vector and the EGR1 promoter.

To prepare 96 well-plates for cell culture, two mls of a coating solution (1:30 dilution of collagen type I (Upstate Biotech Inc. Cat#08-115) in 30% ethanol (filter sterilized)) is added per one 10 cm plate or 50 ml per well of the 96-well plate, and allowed to air dry for 2 hr.

PC12 cells are routinely grown in RPMI-1640 medium (Bio Whittaker) containing 10% horse serum (JRH BIOSCIENCES, Cat. # 12449-78P), 5% heat-inactivated fetal bovine serum (FBS) supplemented with 100 units/ml penicillin and 100 ug/ml streptomycin on a precoated 10 cm tissue culture dish. One to four split is done every three to four days. Cells

are removed from the plates by scraping and resuspended with pipetting up and down for more than 15 times.

Transfect the EGR/SEAP/Neo construct into PC12 using the Lipofectamine protocol described in Example 31. EGR-SEAP/PC12 stable cells are obtained by growing the cells in
5 300 ug/ml G418. The G418-free medium is used for routine growth but every one to two months, the cells should be re-grown in 300 ug/ml G418 for couple of passages.

To assay for neuronal activity, a 10 cm plate with cells around 70 to 80% confluent is screened by removing the old medium. Wash the cells once with PBS (Phosphate buffered saline). Then starve the cells in low serum medium (RPMI-1640 containing 1% horse serum
10 and 0.5% FBS with antibiotics) overnight.

The next morning, remove the medium and wash the cells with PBS. Scrape off the cells from the plate, suspend the cells well in 2 ml low serum medium. Count the cell number and add more low serum medium to reach final cell density as 5×10^5 cells/ml.

Add 200 ul of the cell suspension to each well of 96-well plate (equivalent to 1×10^5
15 cells/well). Add 50 ul supernatant produced by Example 31, 37 degree C for 48 to 72 hr. As a positive control, a growth factor known to activate PC12 cells through EGR can be used, such as 50 ng/ul of Neuronal Growth Factor (NGF). Over fifty-fold induction of SEAP is typically seen in the positive control wells. SEAP assay the supernatant according to Example 37.

20

Example 36: High-Throughput Screening Assay for T-cell Activity

NF-KB (Nuclear Factor KB) is a transcription factor activated by a wide variety of agents including the inflammatory cytokines IL-1 and TNF, CD30 and CD40, lymphotoxin-alpha and lymphotoxin-beta, by exposure to LPS or thrombin, and by expression of certain
25 viral gene products. As a transcription factor, NF-KB regulates the expression of genes involved in immune cell activation, control of apoptosis (NF- KB appears to shield cells from apoptosis), B and T-cell development, anti-viral and antimicrobial responses, and multiple stress responses.

In non-stimulated conditions, NF- KB is retained in the cytoplasm with I-KB
30 (Inhibitor KB). However, upon stimulation, I- KB is phosphorylated and degraded, causing NF- KB to shuttle to the nucleus, thereby activating transcription of target genes. Target genes activated by NF- KB include IL-2, IL-6, GM-CSF, ICAM-1 and class 1 MHC.

Due to its central role and ability to respond to a range of stimuli, reporter constructs utilizing the NF-KB promoter element are used to screen the supernatants produced in Example 31. Activators or inhibitors of NF-KB would be useful in treating, preventing, and/or diagnosing diseases. For example, inhibitors of NF-KB could be used to treat those diseases related to the acute or chronic activation of NF-KB, such as rheumatoid arthritis.

To construct a vector containing the NF-KB promoter element, a PCR based strategy is employed. The upstream primer contains four tandem copies of the NF-KB binding site (GGGGACTTTCCC) (SEQ ID NO:8), 18 bp of sequence complementary to the 5' end of the SV40 early promoter sequence, and is flanked with an XhoI site:

10 5':GCGGCCTCGAGGGGACTTTCCCGGGGACTTTCCGGGGACTTTCCGGGACTTTC
CATCCTGCCATCTCAATTAG:3' (SEQ ID NO:9)

The downstream primer is complementary to the 3' end of the SV40 promoter and is flanked with a Hind III site:

5':GCGGCAAGCTTTTTGCAAAGCCTAGGC:3' (SEQ ID NO:4)

15 PCR amplification is performed using the SV40 promoter template present in the pB-gal:promoter plasmid obtained from Clontech. The resulting PCR fragment is digested with XhoI and Hind III and subcloned into BLSK2-. (Stratagene) Sequencing with the T7 and T3 primers confirms the insert contains the following sequence:

5':CTCGAGGGGACTTTCCCGGGGACTTTCCGGGGACTTTCCGGGACTTTCCATCTG
20 CCATCTCAATTAGTCAGCAACCATAGTCCCGCCCCTAACTCCGCCCATCCCGCCC
CTAACTCCGCCCAGTTCCGCCCATTCTCCGCCCCATGGCTGACTAATTTTTTTTAT
TTATGCAGAGGCCGAGGCCGCCTCGGCCTCTGAGCTATTCCAGAAGTAGTGAGG
AGGCTTTTTTTGGAGGCCTAGGCTTTTGCAAAAAGCTT:3' (SEQ ID NO:10)

Next, replace the SV40 minimal promoter element present in the pSEAP2-promoter plasmid (Clontech) with this NF-KB/SV40 fragment using XhoI and HindIII. However, this vector does not contain a neomycin resistance gene, and therefore, is not preferred for mammalian expression systems.

In order to generate stable mammalian cell lines, the NF-KB/SV40/SEAP cassette is removed from the above NF-KB/SEAP vector using restriction enzymes SalI and NotI, and inserted into a vector containing neomycin resistance. Particularly, the NF-KB/SV40/SEAP cassette was inserted into pGFP-1 (Clontech), replacing the GFP gene, after restricting pGFP-1 with SalI and NotI.

Once NF-KB/SV40/SEAP/Neo vector is created, stable Jurkat T-cells are created and maintained according to the protocol described in Example 33. Similarly, the method for assaying supernatants with these stable Jurkat T-cells is also described in Example 33. As a positive control, exogenous TNF alpha (0.1, 1, 10 ng) is added to wells H9, H10, and H11, with a 5-10 fold activation typically observed.

Example 37: Assay for SEAP Activity

As a reporter molecule for the assays described in Examples 33-36, SEAP activity is assayed using the Tropix Phospho-light Kit (Cat. BP-400) according to the following general procedure. The Tropix Phospho-light Kit supplies the Dilution, Assay, and Reaction Buffers used below.

Prime a dispenser with the 2.5x Dilution Buffer and dispense 15 ul of 2.5x dilution buffer into Optiplates containing 35 ul of a supernatant. Seal the plates with a plastic sealer and incubate at 65 degree C for 30 min. Separate the Optiplates to avoid uneven heating.

Cool the samples to room temperature for 15 minutes. Empty the dispenser and prime with the Assay Buffer. Add 50 ml Assay Buffer and incubate at room temperature 5 min. Empty the dispenser and prime with the Reaction Buffer (see the table below). Add 50 ul Reaction Buffer and incubate at room temperature for 20 minutes. Since the intensity of the chemiluminescent signal is time dependent, and it takes about 10 minutes to read 5 plates on luminometer, one should treat 5 plates at each time and start the second set 10 minutes later.

Read the relative light unit in the luminometer. Set H12 as blank, and print the results. An increase in chemiluminescence indicates reporter activity.

Reaction Buffer Formulation:

# of plates	Rxn buffer diluent (ml)	CSPD (ml)
10	60	3
11	65	3.25
12	70	3.5
13	75	3.75
14	80	4
15	85	4.25
16	90	4.5
17	95	4.75

18	100	5
19	105	5.25
20	110	5.5
21	115	5.75
22	120	6
23	125	6.25
24	130	6.5
25	135	6.75
26	140	7
27	145	7.25
28	150	7.5
29	155	7.75
30	160	8
31	165	8.25
32	170	8.5
33	175	8.75
34	180	9
35	185	9.25
36	190	9.5
37	195	9.75
38	200	10
39	205	10.25
40	210	10.5
41	215	10.75
42	220	11
43	225	11.25
44	230	11.5
45	235	11.75
46	240	12
47	245	12.25
48	250	12.5
49	255	12.75
50	260	13

Example 38: High-Throughput Screening Assay Identifying Changes in Small Molecule Concentration and Membrane Permeability

Binding of a ligand to a receptor is known to alter intracellular levels of small molecules, such as calcium, potassium, sodium, and pH, as well as alter membrane potential. These alterations can be measured in an assay to identify supernatants which bind to receptors of a particular cell. Although the following protocol describes an assay for calcium, this protocol can easily be modified to detect changes in potassium, sodium, pH, membrane potential, or any other small molecule which is detectable by a fluorescent probe.

The following assay uses Fluorometric Imaging Plate Reader ("FLIPR") to measure changes in fluorescent molecules (Molecular Probes) that bind small molecules. Clearly, any fluorescent molecule detecting a small molecule can be used instead of the calcium fluorescent molecule, fluo-4 (Molecular Probes, Inc.; catalog no. F-14202), used here.

For adherent cells, seed the cells at 10,000 -20,000 cells/well in a Co-star black 96-well plate with clear bottom. The plate is incubated in a CO₂ incubator for 20 hours. The adherent cells are washed two times in Biotek washer with 200 ul of HBSS (Hank's Balanced Salt Solution) leaving 100 ul of buffer after the final wash.

A stock solution of 1 mg/ml fluo-4 is made in 10% pluronic acid DMSO. To load the cells with fluo-4, 50 ul of 12 ug/ml fluo-4 is added to each well. The plate is incubated at 37 degrees C in a CO₂ incubator for 60 min. The plate is washed four times in the Biotek washer with HBSS leaving 100 ul of buffer.

For non-adherent cells, the cells are spun down from culture media. Cells are resuspended to $2-5 \times 10^6$ cells/ml with HBSS in a 50-ml conical tube. 4 ul of 1 mg/ml fluo-4 solution in 10% pluronic acid DMSO is added to each ml of cell suspension. The tube is then placed in a 37 degrees C water bath for 30-60 min. The cells are washed twice with HBSS, resuspended to 1×10^6 cells/ml, and dispensed into a microplate, 100 ul/well. The plate is centrifuged at 1000 rpm for 5 min. The plate is then washed once in Denley Cell Wash with 200 ul, followed by an aspiration step to 100 ul final volume.

For a non-cell based assay, each well contains a fluorescent molecule, such as fluo-4. The supernatant is added to the well, and a change in fluorescence is detected.

To measure the fluorescence of intracellular calcium, the FLIPR is set for the following parameters: (1) System gain is 300-800 mW; (2) Exposure time is 0.4 second; (3) Camera F/stop is F/2; (4) Excitation is 488 nm; (5) Emission is 530 nm; and (6) Sample

addition is 50 ul. Increased emission at 530 nm indicates an extracellular signaling event caused by the a molecule, either polypeptide of the present invention or a molecule induced by polypeptide of the present invention, which has resulted in an increase in the intracellular Ca^{++} concentration.

5

Example 40: High-Throughput Screening Assay Identifying Tyrosine Kinase Activity

The Protein Tyrosine Kinases (PTK) represent a diverse group of transmembrane and cytoplasmic kinases. Within the Receptor Protein Tyrosine Kinase (RPTK) group are receptors for a range of mitogenic and metabolic growth factors including the PDGF, FGF, EGF, NGF, HGF and Insulin receptor subfamilies. In addition there are a large family of RPTKs for which the corresponding ligand is unknown. Ligands for RPTKs include mainly secreted small proteins, but also membrane-bound and extracellular matrix proteins.

Activation of RPTK by ligands involves ligand-mediated receptor dimerization, resulting in transphosphorylation of the receptor subunits and activation of the cytoplasmic tyrosine kinases. The cytoplasmic tyrosine kinases include receptor associated tyrosine kinases of the src-family (e.g., src, yes, lck, lyn, fyn) and non-receptor linked and cytosolic protein tyrosine kinases, such as the Jak family, members of which mediate signal transduction triggered by the cytokine superfamily of receptors (e.g., the Interleukins, Interferons, GM-CSF, and Leptin).

Because of the wide range of known factors capable of stimulating tyrosine kinase activity, identifying whether polypeptide of the present invention or a molecule induced by polypeptide of the present invention is capable of activating tyrosine kinase signal transduction pathways is of interest. Therefore, the following protocol is designed to identify such molecules capable of activating the tyrosine kinase signal transduction pathways.

Seed target cells (e.g., primary keratinocytes) at a density of approximately 25,000 cells per well in a 96 well Loprodyne Silent Screen Plates purchased from Nalge Nunc (Naperville, IL). The plates are sterilized with two 30 minute rinses with 100% ethanol, rinsed with water and dried overnight. Some plates are coated for 2 hr with 100 ml of cell culture grade type I collagen (50 mg/ml), gelatin (2%) or polylysine (50 mg/ml), all of which can be purchased from Sigma Chemicals (St. Louis, MO) or 10% Matrigel purchased from Becton Dickinson (Bedford, MA), or calf serum, rinsed with PBS and stored at 4 degree C. Cell growth on these plates is assayed by seeding 5,000 cells/well in growth medium and

indirect quantitation of cell number through use of alamarBlue as described by the manufacturer Alamar Biosciences, Inc. (Sacramento, CA) after 48 hr. Falcon plate covers #3071 from Becton Dickinson (Bedford, MA) are used to cover the Loprodyne Silent Screen Plates. Falcon Microtest III cell culture plates can also be used in some proliferation experiments.

To prepare extracts, A431 cells are seeded onto the nylon membranes of Loprodyne plates (20,000/200ml/well) and cultured overnight in complete medium. Cells are quiesced by incubation in serum-free basal medium for 24 hr. After 5-20 minutes treatment with EGF (60ng/ml) or 50 ul of the supernatant produced in Example 31, the medium was removed and 100 ml of extraction buffer ((20 mM HEPES pH 7.5, 0.15 M NaCl, 1% Triton X-100, 0.1% SDS, 2 mM Na₃VO₄, 2 mM Na₄P₂O₇ and a cocktail of protease inhibitors (# 1836170) obtained from Boehringer Mannheim (Indianapolis, IN) is added to each well and the plate is shaken on a rotating shaker for 5 minutes at 4°C. The plate is then placed in a vacuum transfer manifold and the extract filtered through the 0.45 mm membrane bottoms of each well using house vacuum. Extracts are collected in a 96-well catch/assay plate in the bottom of the vacuum manifold and immediately placed on ice. To obtain extracts clarified by centrifugation, the content of each well, after detergent solubilization for 5 minutes, is removed and centrifuged for 15 minutes at 4 degree C at 16,000 x g.

Test the filtered extracts for levels of tyrosine kinase activity. Although many methods of detecting tyrosine kinase activity are known, one method is described here.

Generally, the tyrosine kinase activity of a supernatant is evaluated by determining its ability to phosphorylate a tyrosine residue on a specific substrate (a biotinylated peptide). Biotinylated peptides that can be used for this purpose include PSK1 (corresponding to amino acids 6-20 of the cell division kinase cdc2-p34) and PSK2 (corresponding to amino acids 1-17 of gastrin). Both peptides are substrates for a range of tyrosine kinases and are available from Boehringer Mannheim.

The tyrosine kinase reaction is set up by adding the following components in order. First, add 10ul of 5uM Biotinylated Peptide, then 10ul ATP/Mg₂⁺ (5mM ATP/50mM MgCl₂), then 10ul of 5x Assay Buffer (40mM imidazole hydrochloride, pH7.3, 40 mM beta-glycerophosphate, 1mM EGTA, 100mM MgCl₂, 5 mM MnCl₂, 0.5 mg/ml BSA), then 5ul of Sodium Vanadate(1mM), and then 5ul of water. Mix the components gently and preincubate

the reaction mix at 30 degree C for 2 min. Initial the reaction by adding 10ul of the control enzyme or the filtered supernatant.

The tyrosine kinase assay reaction is then terminated by adding 10 ul of 120mm EDTA and place the reactions on ice.

- 5 Tyrosine kinase activity is determined by transferring 50 ul aliquot of reaction mixture to a microtiter plate (MTP) module and incubating at 37 degree C for 20 min. This allows the streptavidin coated 96 well plate to associate with the biotinylated peptide. Wash the MTP module with 300ul/well of PBS four times. Next add 75 ul of anti-phosphotyrosine antibody conjugated to horse radish peroxidase(anti-P-Tyr-POD(0.5u/ml)) to each well and
10 incubate at 37 degree C for one hour. Wash the well as above.

Next add 100ul of peroxidase substrate solution (Boehringer Mannheim) and incubate at room temperature for at least 5 mins (up to 30 min). Measure the absorbance of the sample at 405 nm by using ELISA reader. The level of bound peroxidase activity is quantitated using an ELISA reader and reflects the level of tyrosine kinase activity.

15

Example 41: High-Throughput Screening Assay Identifying Phosphorylation Activity

- As a potential alternative and/or compliment to the assay of protein tyrosine kinase activity described in Example 40, an assay which detects activation (phosphorylation) of major intracellular signal transduction intermediates can also be used. For example, as
20 described below one particular assay can detect tyrosine phosphorylation of the Erk-1 and Erk-2 kinases. However, phosphorylation of other molecules, such as Raf, JNK, p38 MAP, Map kinase kinase (MEK), MEK kinase, Src, Muscle specific kinase (MuSK), IRAK, Tec, and Janus, as well as any other phosphoserine, phosphotyrosine, or phosphothreonine molecule, can be detected by substituting these molecules for Erk-1 or Erk-2 in the following
25 assay.

- Specifically, assay plates are made by coating the wells of a 96-well ELISA plate with 0.1ml of protein G (1ug/ml) for 2 hr at room temp, (RT). The plates are then rinsed with PBS and blocked with 3% BSA/PBS for 1 hr at RT. The protein G plates are then treated with 2 commercial monoclonal antibodies (100ng/well) against Erk-1 and Erk-2 (1 hr at RT) (Santa
30 Cruz Biotechnology). (To detect other molecules, this step can easily be modified by substituting a monoclonal antibody detecting any of the above described molecules.) After 3-5 rinses with PBS, the plates are stored at 4 degree C until use.

A431 cells are seeded at 20,000/well in a 96-well Loprodyne filterplate and cultured overnight in growth medium. The cells are then starved for 48 hr in basal medium (DMEM) and then treated with EGF (6ng/well) or 50 ul of the supernatants obtained in Example 31 for 5-20 minutes. The cells are then solubilized and extracts filtered directly into the assay plate.

5 After incubation with the extract for 1 hr at RT, the wells are again rinsed. As a positive control, a commercial preparation of MAP kinase (10ng/well) is used in place of A431 extract. Plates are then treated with a commercial polyclonal (rabbit) antibody (1ug/ml) which specifically recognizes the phosphorylated epitope of the Erk-1 and Erk-2 kinases (1 hr at RT). This antibody is biotinylated by standard procedures. The bound
10 polyclonal antibody is then quantitated by successive incubations with Europium-streptavidin and Europium fluorescence enhancing reagent in the Wallac DELFIA instrument (time-resolved fluorescence). An increased fluorescent signal over background indicates a phosphorylation by polypeptide of the present invention or a molecule induced by polypeptide of the present invention.

15 ***Example 42: Assay for the Stimulation of Bone Marrow CD34+ Cell Proliferation***

This assay is based on the ability of human CD34+ to proliferate in the presence of hematopoietic growth factors and evaluates the ability of isolated polypeptides expressed in mammalian cells to stimulate proliferation of CD34+ cells.

20 It has been previously shown that most mature precursors will respond to only a single signal. More immature precursors require at least two signals to respond. Therefore, to test the effect of polypeptides on hematopoietic activity of a wide range of progenitor cells, the assay contains a given polypeptide in the presence or absence of other hematopoietic growth factors. Isolated cells are cultured for 5 days in the presence of Stem Cell Factor (SCF) in combination with tested sample. SCF alone has a very limited effect on the
25 proliferation of bone marrow (BM) cells, acting in such conditions only as a "survival" factor. However, combined with any factor exhibiting stimulatory effect on these cells (e.g., IL-3), SCF will cause a synergistic effect. Therefore, if the tested polypeptide has a stimulatory effect on a hematopoietic progenitors, such activity can be easily detected. Since
30 normal BM cells have a low level of cycling cells, it is likely that any inhibitory effect of a given polypeptide, or agonists or antagonists thereof, might not be detected. Accordingly, assays for an inhibitory effect on progenitors is preferably tested in cells that are first

subjected to *in vitro* stimulation with SCF+IL+3, and then contacted with the compound that is being evaluated for inhibition of such induced proliferation.

Briefly, CD34+ cells are isolated using methods known in the art. The cells are thawed and resuspended in medium (QBSF 60 serum-free medium with 1% L-glutamine (500ml) Quality Biological, Inc., Gaithersburg, MD Cat# 160-204-101). After several gentle centrifugation steps at 200 x g, cells are allowed to rest for one hour. The cell count is adjusted to 2.5×10^5 cells/ml. During this time, 100 μ l of sterile water is added to the peripheral wells of a 96-well plate. The cytokines that can be tested with a given polypeptide in this assay is rhSCF (R&D Systems, Minneapolis, MN, Cat# 255-SC) at 50 ng/ml alone and in combination with rhSCF and rhIL-3 (R&D Systems, Minneapolis, MN, Cat# 203-ML) at 30 ng/ml. After one hour, 10 μ l of prepared cytokines, 50 μ l of the supernatants prepared in Example 31 (supernatants at 1:2 dilution = 50 μ l) and 20 μ l of diluted cells are added to the media which is already present in the wells to allow for a final total volume of 100 μ l. The plates are then placed in a 37°C/5% CO₂ incubator for five days.

Eighteen hours before the assay is harvested, 0.5 μ Ci/well of [3H] Thymidine is added in a 10 μ l volume to each well to determine the proliferation rate. The experiment is terminated by harvesting the cells from each 96-well plate to a filtermat using the Tomtec Harvester 96. After harvesting, the filtermats are dried, trimmed and placed into OmniFilter assemblies consisting of one OmniFilter plate and one OmniFilter Tray. 60 μ l Microscint is added to each well and the plate sealed with TopSeal-A press-on sealing film. A bar code sticker is affixed to the first plate for counting. The sealed plates is then loaded and the level of radioactivity determined via the Packard Top Count and the printed data collected for analysis. The level of radioactivity reflects the amount of cell proliferation.

The studies described in this example test the activity of a given polypeptide to stimulate bone marrow CD34+ cell proliferation. One skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides (e.g., gene therapy), antibodies, agonists, and/or antagonists and fragments and variants thereof. As a nonlimiting example, potential antagonists tested in this assay would be expected to inhibit cell proliferation in the presence of cytokines and/or to increase the inhibition of cell proliferation in the presence of cytokines and a given polypeptide. In contrast, potential agonists tested in this assay would

be expected to enhance cell proliferation and/or to decrease the inhibition of cell proliferation in the presence of cytokines and a given polypeptide.

The ability of a gene to stimulate the proliferation of bone marrow CD34+ cells indicates that polynucleotides and polypeptides corresponding to the gene are useful for the diagnosis and treatment of disorders affecting the immune system and hematopoiesis. Representative uses are described in the "Immune Activity" and "Infectious Disease" sections above, and elsewhere herein.

Example 43: Assay for Extracellular Matrix Enhanced Cell Response (EMECCR)

The objective of the Extracellular Matrix Enhanced Cell Response (EMECCR) assay is to identify gene products (e.g., isolated polypeptides) that act on the hematopoietic stem cells in the context of the extracellular matrix (ECM) induced signal.

Cells respond to the regulatory factors in the context of signal(s) received from the surrounding microenvironment. For example, fibroblasts, and endothelial and epithelial stem cells fail to replicate in the absence of signals from the ECM. Hematopoietic stem cells can undergo self-renewal in the bone marrow, but not in *in vitro* suspension culture. The ability of stem cells to undergo self-renewal *in vitro* is dependent upon their interaction with the stromal cells and the ECM protein fibronectin (fn). Adhesion of cells to fn is mediated by the $\alpha_5\beta_1$ and $\alpha_4\beta_1$ integrin receptors, which are expressed by human and mouse hematopoietic stem cells. The factor(s) which integrate with the ECM environment and responsible for stimulating stem cell self-renewal has not yet been identified. Discovery of such factors should be of great interest in gene therapy and bone marrow transplant applications

Briefly, polystyrene, non tissue culture treated, 96-well plates are coated with fn fragment at a coating concentration of $0.2 \mu\text{g}/\text{cm}^2$. Mouse bone marrow cells are plated (1,000 cells/well) in 0.2 ml of serum-free medium. Cells cultured in the presence of IL-3 (5 ng/ml) + SCF (50 ng/ml) would serve as the positive control, conditions under which little self-renewal but pronounced differentiation of the stem cells is to be expected. Gene products of the invention (e.g., including, but not limited to, polynucleotides and polypeptides of the present invention, and supernatants produced in Example 31), are tested with appropriate negative controls in the presence and absence of SCF(5.0 ng/ml), where test factor supernates represent 10% of the total assay volume. The plated cells are then allowed to grow by incubating in a low oxygen environment (5% CO₂, 7% O₂, and 88% N₂) tissue

culture incubator for 7 days. The number of proliferating cells within the wells is then quantitated by measuring thymidine incorporation into cellular DNA. Verification of the positive hits in the assay will require phenotypic characterization of the cells, which can be accomplished by scaling up of the culture system and using appropriate antibody reagents against cell surface antigens and FACScan.

One skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides (e.g., gene therapy), antibodies, agonists, and/or antagonists and fragments and variants thereof.

If a particular polypeptide of the present invention is found to be a stimulator of hematopoietic progenitors, polynucleotides and polypeptides corresponding to the gene encoding said polypeptide may be useful for the diagnosis and treatment of disorders affecting the immune system and hematopoiesis. Representative uses are described in the "Immune Activity" and "Infectious Disease" sections above, and elsewhere herein. The gene product may also be useful in the expansion of stem cells and committed progenitors of various blood lineages, and in the differentiation and/or proliferation of various cell types.

Additionally, the polynucleotides and/or polypeptides of the gene of interest and/or agonists and/or antagonists thereof, may also be employed to inhibit the proliferation and differentiation of hematopoietic cells and therefore may be employed to protect bone marrow stem cells from chemotherapeutic agents during chemotherapy. This antiproliferative effect may allow administration of higher doses of chemotherapeutic agents and, therefore, more effective chemotherapeutic treatment.

Moreover, polynucleotides and polypeptides corresponding to the gene of interest may also be useful for the treatment and diagnosis of hematopoietic related disorders such as, for example, anemia, pancytopenia, leukopenia, thrombocytopenia or leukemia since stromal cells are important in the production of cells of hematopoietic lineages. The uses include bone marrow cell ex-vivo culture, bone marrow transplantation, bone marrow reconstitution, radiotherapy or chemotherapy of neoplasia.

Example 44: Human Dermal Fibroblast and Aortic Smooth Muscle Cell Proliferation

The polypeptide of interest is added to cultures of normal human dermal fibroblasts (NHDF) and human aortic smooth muscle cells (AoSMC) and two co-assays are performed with each sample. The first assay examines the effect of the polypeptide of interest on the

proliferation of normal human dermal fibroblasts (NHDF) or aortic smooth muscle cells (AoSMC). Aberrant growth of fibroblasts or smooth muscle cells is a part of several pathological processes, including fibrosis, and restenosis. The second assay examines IL6 production by both NHDF and SMC. IL6 production is an indication of functional activation.

- 5 Activated cells will have increased production of a number of cytokines and other factors, which can result in a proinflammatory or immunomodulatory outcome. Assays are run with and without co-TNF α stimulation, in order to check for costimulatory or inhibitory activity.

Briefly, on day 1, 96-well black plates are set up with 1000 cells/well (NHDF) or 2000 cells/well (AoSMC) in 100 μ l culture media. NHDF culture media contains: Clonetics
10 FB basal media, 1mg/ml hFGF, 5mg/ml insulin, 50mg/ml gentamycin, 2%FBS, while AoSMC culture media contains Clonetics SM basal media, 0.5 μ g/ml hEGF, 5mg/ml insulin, 1 μ g/ml hFGF, 50mg/ml gentamycin, 50 μ g/ml Amphotericin B, 5%FBS. After incubation at 37°C for at least 4-5 hours culture media is aspirated and replaced with growth arrest media. Growth arrest media for NHDF contains fibroblast basal media, 50mg/ml gentamycin, 2%
15 FBS, while growth arrest media for AoSMC contains SM basal media, 50mg/ml gentamycin, 50 μ g/ml Amphotericin B, 0.4% FBS. Incubate at 37°C until day 2.

On day 2, serial dilutions and templates of the polypeptide of interest are designed such that they always include media controls and known-protein controls. For both stimulation and inhibition experiments, proteins are diluted in growth arrest media. For
20 inhibition experiments, TNF α is added to a final concentration of 2ng/ml (NHDF) or 5ng/ml (AoSMC). Add 1/3 vol media containing controls or polypeptides of the present invention and incubate at 37°C/5% CO $_2$ until day 5.

Transfer 60 μ l from each well to another labeled 96-well plate, cover with a plate-sealer, and store at 4°C until Day 6 (for IL6 ELISA). To the remaining 100 μ l in the cell
25 culture plate, aseptically add Alamar Blue in an amount equal to 10% of the culture volume (10 μ l). Return plates to incubator for 3 to 4 hours. Then measure fluorescence with excitation at 530nm and emission at 590nm using the CytoFluor. This yields the growth stimulation/inhibition data.

On day 5, the IL6 ELISA is performed by coating a 96 well plate with 50-100 μ l/well
30 of Anti-Human IL6 Monoclonal antibody diluted in PBS, pH 7.4, incubate ON at room temperature.

On day 6, empty the plates into the sink and blot on paper towels. Prepare Assay Buffer containing PBS with 4% BSA. Block the plates with 200 μ l/well of Pierce Super Block blocking buffer in PBS for 1-2 hr and then wash plates with wash buffer (PBS, 0.05% Tween-20). Blot plates on paper towels. Then add 50 μ l/well of diluted Anti-Human IL-6
5 Monoclonal, Biotin-labeled antibody at 0.50 mg/ml. Make dilutions of IL-6 stock in media (30, 10, 3, 1, 0.3, 0 ng/ml). Add duplicate samples to top row of plate. Cover the plates and incubate for 2 hours at RT on shaker.

Plates are washed with wash buffer and blotted on paper towels. Dilute EU-labeled Streptavidin 1:1000 in Assay buffer, and add 100 μ l/well. Cover the plate and incubate 1 h at
10 RT. Plates are again washed with wash buffer and blotted on paper towels.

Add 100 μ l/well of Enhancement Solution. Shake for 5 minutes. Read the plate on the Wallac DELFIA Fluorometer. Readings from triplicate samples in each assay were tabulated and averaged.

A positive result in this assay suggests AoSMC cell proliferation and that the
15 polypeptide of the present invention may be involved in dermal fibroblast proliferation and/or smooth muscle cell proliferation. A positive result also suggests many potential uses of polypeptides, polynucleotides, agonists and/or antagonists of the polynucleotide/polypeptide of the present invention which gives a positive result. For example, inflammation and immune responses, wound healing, and angiogenesis, as detailed throughout this
20 specification. Particularly, polypeptides of the present invention and polynucleotides of the present invention may be used in wound healing and dermal regeneration, as well as the promotion of vasculargenesis, both of the blood vessels and lymphatics. The growth of vessels can be used in the treatment of, for example, cardiovascular diseases. Additionally, antagonists of polypeptides and polynucleotides of the invention may be useful in treating
25 diseases, disorders, and/or conditions which involve angiogenesis by acting as an anti-vascular (e.g., anti-angiogenesis). These diseases, disorders, and/or conditions are known in the art and/or are described herein, such as, for example, malignancies, solid tumors, benign tumors, for example hemangiomas, acoustic neuromas, neurofibromas, trachomas, and pyogenic granulomas; arteriosclerotic plaques; ocular angiogenic diseases, for example,
30 diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, retinoblastoma, uveitis and Pterygia (abnormal blood vessel growth) of the eye; rheumatoid arthritis; psoriasis; delayed

wound healing; endometriosis; vasculogenesis; granulations; hypertrophic scars (keloids); nonunion fractures; scleroderma; trachoma; vascular adhesions; myocardial angiogenesis; coronary collaterals; cerebral collaterals; arteriovenous malformations; ischemic limb angiogenesis; Osler-Webber Syndrome; plaque neovascularization; telangiectasia; hemophiliac joints; angiofibroma; fibromuscular dysplasia; wound granulation; Crohn's disease; and atherosclerosis. Moreover, antagonists of polypeptides and polynucleotides of the invention may be useful in treating anti-hyperproliferative diseases and/or anti-inflammatory known in the art and/or described herein.

One skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides (e.g., gene therapy), antibodies, agonists, and/or antagonists and fragments and variants thereof.

Example 45: Cellular Adhesion Molecule (CAM) Expression on Endothelial Cells

The recruitment of lymphocytes to areas of inflammation and angiogenesis involves specific receptor-ligand interactions between cell surface adhesion molecules (CAMs) on lymphocytes and the vascular endothelium. The adhesion process, in both normal and pathological settings, follows a multi-step cascade that involves intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion molecule-1 (VCAM-1), and endothelial leukocyte adhesion molecule-1 (E-selectin) expression on endothelial cells (EC). The expression of these molecules and others on the vascular endothelium determines the efficiency with which leukocytes may adhere to the local vasculature and extravasate into the local tissue during the development of an inflammatory response. The local concentration of cytokines and growth factor participate in the modulation of the expression of these CAMs.

Briefly, endothelial cells (e.g., Human Umbilical Vein Endothelial cells (HUVECs)) are grown in a standard 96 well plate to confluence, growth medium is removed from the cells and replaced with 100 μ l of 199 Medium (10% fetal bovine serum (FBS)). Samples for testing and positive or negative controls are added to the plate in triplicate (in 10 μ l volumes). Plates are then incubated at 37°C for either 5 h (selectin and integrin expression) or 24 h (integrin expression only). Plates are aspirated to remove medium and 100 μ l of 0.1% paraformaldehyde-PBS(with Ca⁺⁺ and Mg⁺⁺) is added to each well. Plates are held at 4°C for 30 min. Fixative is removed from the wells and wells are washed 1X with PBS(+Ca,Mg) + 0.5% BSA and drained. 10 μ l of diluted primary antibody is added to the test and control

wells. Anti-ICAM-1-Biotin, Anti-VCAM-1-Biotin and Anti-E-selectin-Biotin are used at a concentration of 10 µg/ml (1:10 dilution of 0.1 mg/ml stock antibody). Cells are incubated at 37°C for 30 min. in a humidified environment. Wells are washed three times with PBS(+Ca,Mg) + 0.5% BSA. 20 µl of diluted ExtrAvidin-Alkaline Phosphatase (1:5,000 dilution, referred to herein as the working dilution) are added to each well and incubated at 37°C for 30 min. Wells are washed three times with PBS(+Ca,Mg)+0.5% BSA. Dissolve 1 tablet of p-Nitrophenol Phosphate pNPP per 5 ml of glycine buffer (pH 10.4). 100 µl of pNPP substrate in glycine buffer is added to each test well. Standard wells in triplicate are prepared from the working dilution of the ExtrAvidin-Alkaline Phosphatase in glycine buffer: 1:5,000 (10^0) > $10^{-0.5}$ > 10^{-1} > $10^{-1.5}$. 5 µl of each dilution is added to triplicate wells and the resulting AP content in each well is 5.50 ng, 1.74 ng, 0.55 ng, 0.18 ng. 100 µl of pNPP reagent is then added to each of the standard wells. The plate is incubated at 37°C for 4h. A volume of 50 µl of 3M NaOH is added to all wells. The plate is read on a plate reader at 405 nm using the background subtraction option on blank wells filled with glycine buffer only. Additionally, the template is set up to indicate the concentration of AP-conjugate in each standard well [5.50 ng; 1.74 ng; 0.55 ng; 0.18 ng]. Results are indicated as amount of bound AP-conjugate in each sample.

Example 46: Alamar Blue Endothelial Cells Proliferation Assay

This assay may be used to quantitatively determine protein mediated inhibition of bFGF-induced proliferation of Bovine Lymphatic Endothelial Cells (LECs), Bovine Aortic Endothelial Cells (BAECs) or Human Microvascular Uterine Myometrial Cells (UTMECs). This assay incorporates a fluorometric growth indicator based on detection of metabolic activity. A standard Alamar Blue Proliferation Assay is prepared in EGM-2MV with 10 ng /ml of bFGF added as a source of endothelial cell stimulation. This assay may be used with a variety of endothelial cells with slight changes in growth medium and cell concentration. Dilutions of the protein batches to be tested are diluted as appropriate. Serum-free medium (GIBCO SFM) without bFGF is used as a non-stimulated control and Angiostatin or TSP-1 are included as a known inhibitory controls.

Briefly, LEC, BAECs or UTMECs are seeded in growth media at a density of 5000 to 2000 cells/well in a 96 well plate and placed at 37-C overnight. After the overnight incubation of the cells, the growth media is removed and replaced with GIBCO EC-SFM.

The cells are treated with the appropriate dilutions of the protein of interest or control protein sample(s) (prepared in SFM) in triplicate wells with additional bFGF to a concentration of 10 ng/ ml. Once the cells have been treated with the samples, the plate(s) is/are placed back in the 37° C incubator for three days. After three days 10 ml of stock alamar blue (Biosource Cat# DAL1100) is added to each well and the plate(s) is/are placed back in the 37° C incubator for four hours. The plate(s) are then read at 530nm excitation and 590nm emission using the CytoFluor fluorescence reader. Direct output is recorded in relative fluorescence units.

Alamar blue is an oxidation-reduction indicator that both fluoresces and changes color in response to chemical reduction of growth medium resulting from cell growth. As cells grow in culture, innate metabolic activity results in a chemical reduction of the immediate surrounding environment. Reduction related to growth causes the indicator to change from oxidized (non-fluorescent blue) form to reduced (fluorescent red) form. i.e. stimulated proliferation will produce a stronger signal and inhibited proliferation will produce a weaker signal and the total signal is proportional to the total number of cells as well as their metabolic activity. The background level of activity is observed with the starvation medium alone. This is compared to the output observed from the positive control samples (bFGF in growth medium) and protein dilutions.

Example 47: Detection of Inhibition of a Mixed Lymphocyte Reaction

This assay can be used to detect and evaluate inhibition of a Mixed Lymphocyte Reaction (MLR) by gene products (e.g., isolated polypeptides). Inhibition of a MLR may be due to a direct effect on cell proliferation and viability, modulation of costimulatory molecules on interacting cells, modulation of adhesiveness between lymphocytes and accessory cells, or modulation of cytokine production by accessory cells. Multiple cells may be targeted by these polypeptides since the peripheral blood mononuclear fraction used in this assay includes T, B and natural killer lymphocytes, as well as monocytes and dendritic cells.

Polypeptides of interest found to inhibit the MLR may find application in diseases associated with lymphocyte and monocyte activation or proliferation. These include, but are not limited to, diseases such as asthma, arthritis, diabetes, inflammatory skin conditions, psoriasis, eczema, systemic lupus erythematosus, multiple sclerosis, glomerulonephritis, inflammatory bowel disease, crohn's disease, ulcerative colitis, arteriosclerosis, cirrhosis,

graft vs. host disease, host vs. graft disease, hepatitis, leukemia and lymphoma.

Briefly, PBMCs from human donors are purified by density gradient centrifugation using Lymphocyte Separation Medium (LSM[®], density 1.0770 g/ml, Organon Teknika Corporation, West Chester, PA). PBMCs from two donors are adjusted to 2×10^6 cells/ml in
5 RPMI-1640 (Life Technologies, Grand Island, NY) supplemented with 10% FCS and 2 mM glutamine. PBMCs from a third donor is adjusted to 2×10^5 cells/ml. Fifty microliters of PBMCs from each donor is added to wells of a 96-well round bottom microtiter plate. Dilutions of test materials (50 μ l) is added in triplicate to microtiter wells. Test samples (of the protein of interest) are added for final dilution of 1:4; rhuIL-2 (R&D Systems,
10 Minneapolis, MN, catalog number 202-IL) is added to a final concentration of 1 μ g/ml; anti-CD4 mAb (R&D Systems, clone 34930.11, catalog number MAB379) is added to a final concentration of 10 μ g/ml. Cells are cultured for 7-8 days at 37°C in 5% CO₂, and 1 μ C of [³H] thymidine is added to wells for the last 16 hrs of culture. Cells are harvested and thymidine incorporation determined using a Packard TopCount. Data is expressed as the
15 mean and standard deviation of triplicate determinations.

Samples of the protein of interest are screened in separate experiments and compared to the negative control treatment, anti-CD4 mAb, which inhibits proliferation of lymphocytes and the positive control treatment, IL-2 (either as recombinant material or supernatant), which enhances proliferation of lymphocytes.

20 One skilled in the art could easily modify the exemplified studies to test the activity of polynucleotides (e.g., gene therapy), antibodies, agonists, and/or antagonists and fragments and variants thereof.

Example 48: Assays for Protease Activity

25 The following assay may be used to assess protease activity of the polypeptides of the invention.

Gelatin and casein zymography are performed essentially as described (Heusen et al., *Anal. Biochem.*, 102:196-202 (1980); Wilson et al., *Journal of Urology*, 149:653-658 (1993)). Samples are run on 10% polyacrylamide/0.1% SDS gels containing 1% gelatin or casein,
30 soaked in 2.5% triton at room temperature for 1 hour, and in 0.1M glycine, pH 8.3 at 37°C 5 to 16 hours. After staining in amido black areas of proteolysis appear as clear areas against the blue-black background. Trypsin (Sigma T8642) is used as a positive control.

Protease activity is also determined by monitoring the cleavage of n-a-benzoyl-L-arginine ethyl ester (BAEE) (Sigma B-4500. Reactions are set up in (25mMNaPO₄,1mM EDTA, and 1mM BAEE), pH 7.5. Samples are added and the change in adsorbance at 260nm is monitored on the Beckman DU-6 spectrophotometer in the time-drive mode.

5 Trypsin is used as a positive control.

Additional assays based upon the release of acid-soluble peptides from casein or hemoglobin measured as adsorbance at 280 nm or colorimetrically using the Folin method are performed as described in Bergmeyer, et al., *Methods of Enzymatic Analysis*, 5 (1984). Other assays involve the solubilization of chromogenic substrates (Ward, *Applied Science*,
10 251-317 (1983).

Example 49: Identifying Serine Protease Substrate Specificity

Methods known in the art or described herein may be used to determine the substrate specificity of the polypeptides of the present invention having serine protease activity. A
15 preferred method of determining substrate specificity is by the use of positional scanning synthetic combinatorial libraries as described in GB 2 324 529 (incorporated herein in its entirety).

Example 50: Ligand Binding Assays

20 The following assay may be used to assess ligand binding activity of the polypeptides of the invention.

Ligand binding assays provide a direct method for ascertaining receptor pharmacology and are adaptable to a high throughput format. The purified ligand for a polypeptide is radiolabeled to high specific activity (50-2000 Ci/mmol) for binding studies. A
25 determination is then made that the process of radiolabeling does not diminish the activity of the ligand towards its polypeptide. Assay conditions for buffers, ions, pH and other modulators such as nucleotides are optimized to establish a workable signal to noise ratio for both membrane and whole cell polypeptide sources. For these assays, specific polypeptide binding is defined as total associated radioactivity minus the radioactivity measured in the
30 presence of an excess of unlabeled competing ligand. Where possible, more than one competing ligand is used to define residual nonspecific binding.

Example 51: Functional Assay in *Xenopus* Oocytes

Capped RNA transcripts from linearized plasmid templates encoding the polypeptides of the invention are synthesized in vitro with RNA polymerases in accordance with standard procedures. In vitro transcripts are suspended in water at a final concentration of 0.2 mg/ml.

- 5 Ovarian lobes are removed from adult female toads, Stage V defolliculated oocytes are obtained, and RNA transcripts (10 ng/oocyte) are injected in a 50 nl bolus using a microinjection apparatus. Two electrode voltage clamps are used to measure the currents from individual *Xenopus* oocytes in response polypeptides and polypeptide agonist exposure. Recordings are made in Ca²⁺ free Barth's medium at room temperature. The *Xenopus* system
10 can be used to screen known ligands and tissue/cell extracts for activating ligands.

Example 52: Microphysiometric Assays

Activation of a wide variety of secondary messenger systems results in extrusion of small amounts of acid from a cell. The acid formed is largely as a result of the increased
15 metabolic activity required to fuel the intracellular signaling process. The pH changes in the media surrounding the cell are very small but are detectable by the CYTOSENSOR microphysiometer (Molecular Devices Ltd., Menlo Park, Calif.). The CYTOSENSOR is thus capable of detecting the activation of polypeptide which is coupled to an energy utilizing intracellular signaling pathway.

20

Example 53: Extract/Cell Supernatant Screening

A large number of mammalian receptors exist for which there remains, as yet, no cognate activating ligand (agonist). Thus, active ligands for these receptors may not be included within the ligands banks as identified to date. Accordingly, the polypeptides of the
25 invention can also be functionally screened (using calcium, cAMP, microphysiometer, oocyte electrophysiology, etc., functional screens) against tissue extracts to identify its natural ligands. Extracts that produce positive functional responses can be sequentially subfractionated until an activating ligand is isolated identified.

30 Example 54: Calcium and cAMP Functional Assays

Seven transmembrane receptors which are expressed in HEK 293 cells have been shown to be coupled functionally to activation of PLC and calcium mobilization and/or

cAMP stimulation or inhibition. Basal calcium levels in the HEK 293 cells in receptor-transfected or vector control cells were observed to be in the normal, 100 nM to 200 nM, range. HEK 293 cells expressing recombinant receptors are loaded with fura 2 and in a single day >150 selected ligands or tissue/cell extracts are evaluated for agonist induced calcium mobilization. Similarly, HEK 293 cells expressing recombinant receptors are evaluated for the stimulation or inhibition of cAMP production using standard cAMP quantitation assays. Agonists presenting a calcium transient or cAMP fluctuation are tested in vector control cells to determine if the response is unique to the transfected cells expressing receptor.

10 *Example 55: ATP-binding assay*

The following assay may be used to assess ATP-binding activity of polypeptides of the invention.

ATP-binding activity of the polypeptides of the invention may be detected using the ATP-binding assay described in U.S. Patent 5, 858, 719, which is herein incorporated by reference in its entirety. Briefly, ATP-binding to polypeptides of the invention is measured via photoaffinity labeling with 8-azido-ATP in a competition assay. Reaction mixtures containing 1 mg/ml of the ABC transport protein of the present invention are incubated with varying concentrations of ATP, or the non-hydrolyzable ATP analog adenylyl-5'-imidodiphosphate for 10 minutes at 4°C. A mixture of 8-azido-ATP (Sigma Chem. Corp., St. Louis, MO.) plus 8-azido-ATP (γ -³²P-ATP) (5 mCi/μmol, ICN, Irvine CA.) is added to a final concentration of 100 μM and 0.5 ml aliquots are placed in the wells of a porcelain spot plate on ice. The plate is irradiated using a short wave 254 nm UV lamp at a distance of 2.5 cm from the plate for two one-minute intervals with a one-minute cooling interval in between. The reaction is stopped by addition of dithiothreitol to a final concentration of 2mM. The incubations are subjected to SDS-PAGE electrophoresis, dried, and autoradiographed. Protein bands corresponding to the particular polypeptides of the invention are excised, and the radioactivity quantified. A decrease in radioactivity with increasing ATP or adenylyl-5'-imidodiphosphate provides a measure of ATP affinity to the polypeptides.

30

Example 56: Small Molecule Screening

This invention is particularly useful for screening therapeutic compounds by using the

polypeptides of the invention, or binding fragments thereof, in any of a variety of drug screening techniques. The polypeptide or fragment employed in such a test may be affixed to a solid support, expressed on a cell surface, free in solution, or located intracellularly. One method of drug screening utilizes eukaryotic or prokaryotic host cells which are stably transformed with recombinant nucleic acids expressing the polypeptide or fragment. Drugs are screened against such transformed cells in competitive binding assays. One may measure, for example, the formulation of complexes between the agent being tested and polypeptide of the invention.

Thus, the present invention provides methods of screening for drugs or any other agents which affect activities mediated by the polypeptides of the invention. These methods comprise contacting such an agent with a polypeptide of the invention or fragment thereof and assaying for the presence of a complex between the agent and the polypeptide or fragment thereof, by methods well known in the art. In such a competitive binding assay, the agents to screen are typically labeled. Following incubation, free agent is separated from that present in bound form, and the amount of free or uncomplexed label is a measure of the ability of a particular agent to bind to the polypeptides of the invention.

Another technique for drug screening provides high throughput screening for compounds having suitable binding affinity to the polypeptides of the invention, and is described in great detail in European Patent Application 84/03564, published on September 13, 1984, which is herein incorporated by reference in its entirety. Briefly stated, large numbers of different small molecule test compounds are synthesized on a solid substrate, such as plastic pins or some other surface. The test compounds are reacted with polypeptides of the invention and washed. Bound polypeptides are then detected by methods well known in the art. Purified polypeptides are coated directly onto plates for use in the aforementioned drug screening techniques. In addition, non-neutralizing antibodies may be used to capture the peptide and immobilize it on the solid support.

This invention also contemplates the use of competitive drug screening assays in which neutralizing antibodies capable of binding polypeptides of the invention specifically compete with a test compound for binding to the polypeptides or fragments thereof. In this manner, the antibodies are used to detect the presence of any peptide which shares one or more antigenic epitopes with a polypeptide of the invention.

Example 57: Phosphorylation Assay

In order to assay for phosphorylation activity of the polypeptides of the invention, a phosphorylation assay as described in U.S. Patent 5,958,405 (which is herein incorporated by reference) is utilized. Briefly, phosphorylation activity may be measured by phosphorylation of a protein substrate using gamma-labeled ^{32}P -ATP and quantitation of the incorporated radioactivity using a gamma radioisotope counter. The polypeptides of the invention are incubated with the protein substrate, ^{32}P -ATP, and a kinase buffer. The ^{32}P incorporated into the substrate is then separated from free ^{32}P -ATP by electrophoresis, and the incorporated ^{32}P is counted and compared to a negative control. Radioactivity counts above the negative control are indicative of phosphorylation activity of the polypeptides of the invention.

Example 58: Detection of Phosphorylation Activity (Activation) of the Polypeptides of the Invention in the Presence of Polypeptide Ligands

Methods known in the art or described herein may be used to determine the phosphorylation activity of the polypeptides of the invention. A preferred method of determining phosphorylation activity is by the use of the tyrosine phosphorylation assay as described in US 5,817,471 (incorporated herein by reference).

Example 59: Identification Of Signal Transduction Proteins That Interact With Polypeptides Of The Present Invention

The inventive purified polypeptides of the invention are research tools for the identification, characterization and purification of additional signal transduction pathway proteins or receptor proteins. Briefly, labeled receptor PTK polypeptide is useful as a reagent for the purification of molecules with which it interacts. In one embodiment of affinity purification, receptor PTK polypeptide is covalently coupled to a chromatography column. Cell-free extract derived from putative target cells, such as carcinoma tissues, is passed over the column, and molecules with appropriate affinity bind to the receptor PTK polypeptides, or specific phosphotyrosine-recognition domains thereof. The receptor PTK polypeptide interacting protein-complex is recovered from the column, dissociated, and the recovered molecule subjected to N-terminal protein sequencing. This amino acid sequence is then used to identify the captured molecule or to design degenerate oligonucleotide probes for cloning

the relevant gene from an appropriate cDNA library.

Example 60: IL-6 Bioassay

To test the proliferative effects of the polypeptides of the invention, the IL-6 Bioassay as described by Marz *et al.* is utilized (*Proc. Natl. Acad. Sci., U.S.A.*, 95:3251-56 (1998), which is herein incorporated by reference). Briefly, IL-6 dependent B9 murine cells are washed three times in IL-6 free medium and plated at a concentration of 5,000 cells per well in 50 μ l, and 50 μ l of the IL-6-like polypeptide is added. After 68 hrs. at 37°C, the number of viable cells is measured by adding the tetrazolium salt thiazolyl blue (MTT) and incubating for a further 4 hrs. at 37°C. B9 cells are lysed by SDS and optical density is measured at 570 nm. Controls containing IL-6 (positive) and no cytokine (negative) are utilized. Enhanced proliferation in the test sample(s) relative to the negative control is indicative of proliferative effects mediated by polypeptides of the invention.

Example 61: Support of Chicken Embryo Neuron Survival

To test whether sympathetic neuronal cell viability is supported by polypeptides of the invention, the chicken embryo neuronal survival assay of Senaldi *et al* is utilized (*Proc. Natl. Acad. Sci., U.S.A.*, 96:11458-63 (1998), which is herein incorporated by reference). Briefly, motor and sympathetic neurons are isolated from chicken embryos, resuspended in L15 medium (with 10% FCS, glucose, sodium selenite, progesterone, conalbumin, putrescine, and insulin; Life Technologies, Rockville, MD.) and Dulbecco's modified Eagles medium [with 10% FCS, glutamine, penicillin, and 25 mM Hepes buffer (pH 7.2); Life Technologies, Rockville, MD.], respectively, and incubated at 37°C in 5% CO₂ in the presence of different concentrations of the inventive purified IL-6-like polypeptide, as well as a negative control lacking any cytokine. After 3 days, neuron survival is determined by evaluation of cellular morphology, and through the use of the colorimetric assay of Mosmann (Mossmann, T., *J. Immunol. Methods*, 65:55-63 (1983)). Enhanced neuronal cell viability as compared to the controls lacking cytokine is indicative of the ability of the inventive purified IL-6-like polypeptide(s) to enhance the survival of neuronal cells.

Example 62: Assay for Phosphatase Activity

The following assay may be used to assess serine/threonine phosphatase (PTPase) activity of the polypeptides of the invention.

In order to assay for serine/threonine phosphatase (PTPase) activity, assays can be
5 utilized which are widely known to those skilled in the art. For example, the serine/threonine phosphatase (PSPase) activity is measured using a PSPase assay kit from New England Biolabs, Inc. Myelin basic protein (MyBP), a substrate for PSPase, is phosphorylated on serine and threonine residues with cAMP-dependent Protein Kinase in the presence of [³²P]ATP. Protein serine/threonine phosphatase activity is then determined by measuring the
10 release of inorganic phosphate from ³²P-labeled MyBP.

Example 63: Interaction of Serine/Threonine Phosphatases with other Proteins

The polypeptides of the invention with serine/threonine phosphatase activity as determined in Example 62 are research tools for the identification, characterization and
15 purification of additional interacting proteins or receptor proteins, or other signal transduction pathway proteins. Briefly, labeled polypeptide(s) of the invention is useful as a reagent for the purification of molecules with which it interacts. In one embodiment of affinity purification, polypeptide of the invention is covalently coupled to a chromatography column. Cell-free extract derived from putative target cells, such as neural or liver cells, is passed over
20 the column, and molecules with appropriate affinity bind to the polypeptides of the invention. The polypeptides of the invention -complex is recovered from the column, dissociated, and the recovered molecule subjected to N-terminal protein sequencing. This amino acid sequence is then used to identify the captured molecule or to design degenerate oligonucleotide probes for cloning the relevant gene from an appropriate cDNA library.

25

Example 64: Assaying for Heparanase Activity

In order to assay for heparanase activity of the polypeptides of the invention, the heparanase assay described by Vlodavsky et al is utilized (Vlodavsky, I., et al., Nat. Med., 5:793-802 (1999)). Briefly, cell lysates, conditioned media or intact cells (1 x 10⁶ cells per
30 35-mm dish) are incubated for 18 hrs at 37°C, pH 6.2-6.6, with ³⁵S-labeled ECM or soluble ECM derived peak I proteoglycans. The incubation medium is centrifuged and the supernatant is analyzed by gel filtration on a Sepharose CL-6B column (0.9 x 30 cm).

Fractions are eluted with PBS and their radioactivity is measured. Degradation fragments of heparan sulfate side chains are eluted from Sepharose 6B at $0.5 < K_{av} < 0.8$ (peak II). Each experiment is done at least three times. Degradation fragments corresponding to "peak II," as described by Vlodavsky et al., is indicative of the activity of the polypeptides of the invention in cleaving heparan sulfate.

Example 65: Immobilization of biomolecules

This example provides a method for the stabilization of polypeptides of the invention in non-host cell lipid bilayer constructs (see, e.g., Bieri et al., Nature Biotech 17:1105-1108 (1999), hereby incorporated by reference in its entirety herein) which can be adapted for the study of polypeptides of the invention in the various functional assays described above. Briefly, carbohydrate-specific chemistry for biotinylation is used to confine a biotin tag to the extracellular domain of the polypeptides of the invention, thus allowing uniform orientation upon immobilization. A 50uM solution of polypeptides of the invention in washed membranes is incubated with 20 mM NaIO₄ and 1.5 mg/ml (4mM) BACH or 2 mg/ml (7.5mM) biotin-hydrazide for 1 hr at room temperature (reaction volume, 150ul). Then the sample is dialyzed (Pierce Slidealizer Cassett, 10 kDa cutoff; Pierce Chemical Co., Rockford IL) at 4C first for 5 h, exchanging the buffer after each hour, and finally for 12 h against 500 ml buffer R (0.15 M NaCl, 1 mM MgCl₂, 10 mM sodium phosphate, pH7). Just before addition into a cuvette, the sample is diluted 1:5 in buffer ROG50 (Buffer R supplemented with 50 mM octylglucoside).

Example 66: TAQMAN

Quantitative PCR (QPCR). Total RNA from cells in culture are extracted by Trizol separation as recommended by the supplier (LifeTechnologies). (Total RNA is treated with DNase I (Life Technologies) to remove any contaminating genomic DNA before reverse transcription.) Total RNA (50 ng) is used in a one-step, 50ul, RT-QPCR, consisting of Taqman Buffer A (Perkin-Elmer; 50 mM KCl/10 mM Tris, pH 8.3), 5.5 mM MgCl₂, 240 μM each dNTP, 0.4 units RNase inhibitor(Promega), 8%glycerol, 0.012% Tween-20, 0.05% gelatin, 0.3uM primers, 0.1uM probe, 0.025units Amplitaq Gold (Perkin-Elmer) and 2.5 units Superscript II reverse transcriptase (Life Technologies). As a control for genomic

contamination, parallel reactions are setup without reverse transcriptase. The relative abundance of (unknown) and 18S RNAs are assessed by using the Applied Biosystems Prism 7700 Sequence Detection System (Livak, K. J., Flood, S. J., Marmaro, J., Giusti, W. & Deetz, K. (1995) PCR Methods Appl. 4, 357-362). Reactions are carried out at 48°C for 30 min, 5 95°C for 10 min, followed by 40 cycles of 95°C for 15s, 60°C for 1 min. Reactions are performed in triplicate.

Primers (f & r) and FRET probes sets are designed using Primer Express Software (Perkin-Elmer). Probes are labeled at the 5'-end with the reporter dye 6-FAM and on the 3'-end with the quencher dye TAMRA (Biosource International, Camarillo, CA or Perkin-10 Elmer).

It will be clear that the invention may be practiced otherwise than as particularly described in the foregoing description and examples. Numerous modifications and variations of the present invention are possible in light of the above teachings and, therefore, are within 15 the scope of the appended claims.

The entire disclosure of each document cited (including patents, patent applications, journal articles, abstracts, laboratory manuals, books, or other disclosures) in the Background of the Invention, Detailed Description, and Examples is hereby incorporated herein by reference. Further, the hard copy of the sequence listing submitted herewith and the 20 corresponding computer readable form are both incorporated herein by reference in their entireties. Moreover, the hard copy of and the corresponding computer readable form of the Sequence Listing of U.S. Patent Application Serial Nos. 60/144,972; 60/148,681; 60/149,173; 60/158,004; and 60/194,689 are also incorporated herein by reference in their entireties.

25

30

Applicant's or agent's file referencenumber	PA002PCT	International application No.	UNASSIGNED
--	----------	-------------------------------	------------

INDICATIONS RELATING TO A DEPOSITED MICROORGANISM

(PCT Rule 13bis)

A. The indications made below relate to the microorganism referred to in the description on page <u>78</u> , line <u>N/A</u>	
B. IDENTIFICATION OF DEPOSIT Further deposits are identified on an additional sheet <input type="checkbox"/>	
Name of depositary institution American Type Culture Collection	
Address of depositary institution (including postal code and country) 10801 University Boulevard Manassas, Virginia 20110-2209 United States of America	
Date of deposit 23 July 1999	Accession Number PTA-416
C. ADDITIONAL INDICATIONS (leave blank if not applicable) This information is continued on an additional sheet <input type="checkbox"/>	
D. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE (if the indications are not for all designated States)	
Europe In respect to those designations in which a European Patent is sought a sample of the deposited microorganism will be made available until the publication of the mention of the grant of the European patent or until the date on which application has been refused or withdrawn or is deemed to be withdrawn, only by the issue of such a sample to an expert nominated by the person requesting the sample (Rule 28 (4) EPC). Continued on the Attached Pages 2 & 3	
E. SEPARATE FURNISHING OF INDICATIONS (leave blank if not applicable)	
The indications listed below will be submitted to the International Bureau later (specify the general nature of the indications e.g., "Accession Number of Deposit")	

For receiving Office use only	For International Bureau use only
<input type="checkbox"/> This sheet was received with the international application	<input type="checkbox"/> This sheet was received by the International Bureau on:
Authorized officer	Authorized officer

ATCC Deposit No. PTA-416**Page No. 2****CANADA**

The applicant requests that, until either a Canadian patent has been issued on the basis of an application or the application has been refused, or is abandoned and no longer subject to reinstatement, or is withdrawn, the Commissioner of Patents only authorizes the furnishing of a sample of the deposited biological material referred to in the application to an independent expert nominated by the Commissioner, the applicant must, by a written statement, inform the International Bureau accordingly before completion of technical preparations for publication of the international application.

NORWAY

The applicant hereby requests that the application has been laid open to public inspection (by the Norwegian Patent Office), or has been finally decided upon by the Norwegian Patent Office without having been laid open inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Norwegian Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Norwegian Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on the list of recognized experts drawn up by the Norwegian Patent Office or any person approved by the applicant in the individual case.

AUSTRALIA

The applicant hereby gives notice that the furnishing of a sample of a microorganism shall only be effected prior to the grant of a patent, or prior to the lapsing, refusal or withdrawal of the application, to a person who is a skilled addressee without an interest in the invention (Regulation 3.25(3) of the Australian Patents Regulations).

FINLAND

The applicant hereby requests that, until the application has been laid open to public inspection (by the National Board of Patents and Regulations), or has been finally decided upon by the National Board of Patents and Registration without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art.

UNITED KINGDOM

The applicant hereby requests that the furnishing of a sample of a microorganism shall only be made available to an expert. The request to this effect must be filed by the applicant with the International Bureau before the completion of the technical preparations for the international publication of the application.

ATCC Deposit No.: PTA-416**Page N . 3****DENMARK**

The applicant hereby requests that, until the application has been laid open to public inspection (by the Danish Patent Office), or has been finally decided upon by the Danish Patent office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Danish Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Danish Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Danish Patent Office or any person by the applicant in the individual case.

SWEDEN

The applicant hereby requests that, until the application has been laid open to public inspection (by the Swedish Patent Office), or has been finally decided upon by the Swedish Patent Office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the International Bureau before the expiration of 16 months from the priority date (preferably on the Form PCT/RO/134 reproduced in annex Z of Volume I of the PCT Applicant's Guide). If such a request has been filed by the applicant any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Swedish Patent Office or any person approved by a applicant in the individual case.

NETHERLANDS

The applicant hereby requests that until the date of a grant of a Netherlands patent or until the date on which the application is refused or withdrawn or lapsed, the microorganism shall be made available as provided in the 31F(1) of the Patent Rules only by the issue of a sample to an expert. The request to this effect must be furnished by the applicant with the Netherlands Industrial Property Office before the date on which the application is made available to the public under Section 22C or Section 25 of the Patents Act of the Kingdom of the Netherlands, whichever of the two dates occurs earlier.

340

Applicant's or agent's file referencenumber	PA002PCT	International application No.	UNASSIGNED
--	----------	-------------------------------	------------

INDICATIONS RELATING TO A DEPOSITED MICROORGANISM

(PCT Rule 13bis)

A. The indications made below relate to the microorganism referred to in the description on page <u>78</u> , line <u>N/A</u>	
B. IDENTIFICATION OF DEPOSIT Further deposits are identified on an additional sheet <input type="checkbox"/>	
Name of depositary institution American Type Culture Collection	
Address of depositary institution (including postal code and country) 10801 University Boulevard Manassas, Virginia 20110-2209 United States of America	
Date of deposit 11 August 1999	Accession Number PTA-497
C. ADDITIONAL INDICATIONS (leave blank if not applicable) This information is continued on an additional sheet <input type="checkbox"/>	
D. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE (if the indications are not for all designated States)	
Europe In respect to those designations in which a European Patent is sought a sample of the deposited microorganism will be made available until the publication of the mention of the grant of the European patent or until the date on which application has been refused or withdrawn or is deemed to be withdrawn, only by the issue of such a sample to an expert nominated by the person requesting the sample (Rule 28 (4) EPC). Continued on the Attached Pages 2 & 3	
E. SEPARATE FURNISHING OF INDICATIONS (leave blank if not applicable)	
The indications listed below will be submitted to the International Bureau later (specify the general nature of the indications e.g., "Accession Number of Deposit")	

For receiving Office use only	For International Bureau use only
<input type="checkbox"/> This sheet was received with the international application	<input type="checkbox"/> This sheet was received by the International Bureau on:
Authorized officer	Authorized officer

ATCC Deposit No. PTA-497**Page No. 2****CANADA**

The applicant requests that, until either a Canadian patent has been issued on the basis of an application or the application has been refused, or is abandoned and no longer subject to reinstatement, or is withdrawn, the Commissioner of Patents only authorizes the furnishing of a sample of the deposited biological material referred to in the application to an independent expert nominated by the Commissioner, the applicant must, by a written statement, inform the International Bureau accordingly before completion of technical preparations for publication of the international application.

NORWAY

The applicant hereby requests that the application has been laid open to public inspection (by the Norwegian Patent Office), or has been finally decided upon by the Norwegian Patent Office without having been laid open inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Norwegian Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Norwegian Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on the list of recognized experts drawn up by the Norwegian Patent Office or any person approved by the applicant in the individual case.

AUSTRALIA

The applicant hereby gives notice that the furnishing of a sample of a microorganism shall only be effected prior to the grant of a patent, or prior to the lapsing, refusal or withdrawal of the application, to a person who is a skilled addressee without an interest in the invention (Regulation 3.25(3) of the Australian Patents Regulations).

FINLAND

The applicant hereby requests that, until the application has been laid open to public inspection (by the National Board of Patents and Regulations), or has been finally decided upon by the National Board of Patents and Registration without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art.

UNITED KINGDOM

The applicant hereby requests that the furnishing of a sample of a microorganism shall only be made available to an expert. The request to this effect must be filed by the applicant with the International Bureau before the completion of the technical preparations for the international publication of the application.

ATCC Deposit No.: PTA-497**Page No. 3****DENMARK**

The applicant hereby requests that, until the application has been laid open to public inspection (by the Danish Patent Office), or has been finally decided upon by the Danish Patent office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Danish Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Danish Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Danish Patent Office or any person by the applicant in the individual case.

SWEDEN

The applicant hereby requests that, until the application has been laid open to public inspection (by the Swedish Patent Office), or has been finally decided upon by the Swedish Patent Office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the International Bureau before the expiration of 16 months from the priority date (preferably on the Form PCT/RO/134 reproduced in annex Z of Volume I of the PCT Applicant's Guide). If such a request has been filed by the applicant any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Swedish Patent Office or any person approved by a applicant in the individual case.

NETHERLANDS

The applicant hereby requests that until the date of a grant of a Netherlands patent or until the date on which the application is refused or withdrawn or lapsed, the microorganism shall be made available as provided in the 31F(1) of the Patent Rules only by the issue of a sample to an expert. The request to this effect must be furnished by the applicant with the Netherlands Industrial Property Office before the date on which the application is made available to the public under Section 22C or Section 25 of the Patents Act of the Kingdom of the Netherlands, whichever of the two dates occurs earlier.

Applicant's or agent's file reference number	PA002PCT	International application No.	UNASSIGNED
---	----------	-------------------------------	------------

INDICATIONS RELATING TO A DEPOSITED MICROORGANISM

(PCT Rule 13bis)

A. The indications made below relate to the microorganism referred to in the description on page <u>78</u> , line <u>N/A</u>	
B. IDENTIFICATION OF DEPOSIT Further deposits are identified on an additional sheet <input type="checkbox"/>	
Name of depositary institution American Type Culture Collection	
Address of depositary institution (including postal code and country) 10801 University Boulevard Manassas, Virginia 20110-2209 United States of America	
Date of deposit 03 April 2000	Accession Number PTA-1620
C. ADDITIONAL INDICATIONS (leave blank if not applicable) This information is continued on an additional sheet <input type="checkbox"/>	
D. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE (if the indications are not for all designated States)	
Europe In respect to those designations in which a European Patent is sought a sample of the deposited microorganism will be made available until the publication of the mention of the grant of the European patent or until the date on which application has been refused or withdrawn or is deemed to be withdrawn, only by the issue of such a sample to an expert nominated by the person requesting the sample (Rule 28 (4) EPC). Continued on the Attached Pages 2 & 3	
E. SEPARATE FURNISHING OF INDICATIONS (leave blank if not applicable)	
The indications listed below will be submitted to the International Bureau later (specify the general nature of the indications e.g., "Accession Number of Deposit")	

For receiving Office use only	For International Bureau use only
<input type="checkbox"/> This sheet was received with the international application	<input type="checkbox"/> This sheet was received by the International Bureau on:
Authorized officer	Authorized officer

ATCC Deposit N . PTA-1620**Page N . 2****CANADA**

The applicant requests that, until either a Canadian patent has been issued on the basis of an application or the application has been refused, or is abandoned and no longer subject to reinstatement, or is withdrawn, the Commissioner of Patents only authorizes the furnishing of a sample of the deposited biological material referred to in the application to an independent expert nominated by the Commissioner, the applicant must, by a written statement, inform the International Bureau accordingly before completion of technical preparations for publication of the international application.

NORWAY

The applicant hereby requests that the application has been laid open to public inspection (by the Norwegian Patent Office), or has been finally decided upon by the Norwegian Patent Office without having been laid open inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Norwegian Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Norwegian Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on the list of recognized experts drawn up by the Norwegian Patent Office or any person approved by the applicant in the individual case.

AUSTRALIA

The applicant hereby gives notice that the furnishing of a sample of a microorganism shall only be effected prior to the grant of a patent, or prior to the lapsing, refusal or withdrawal of the application, to a person who is a skilled addressee without an interest in the invention (Regulation 3.25(3) of the Australian Patents Regulations).

FINLAND

The applicant hereby requests that, until the application has been laid open to public inspection (by the National Board of Patents and Regulations), or has been finally decided upon by the National Board of Patents and Registration without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art.

UNITED KINGDOM

The applicant hereby requests that the furnishing of a sample of a microorganism shall only be made available to an expert. The request to this effect must be filed by the applicant with the International Bureau before the completion of the technical preparations for the international publication of the application.

ATCC Dep sit N .: PTA-1620

Page N . 3

DENMARK

The applicant hereby requests that, until the application has been laid open to public inspection (by the Danish Patent Office), or has been finally decided upon by the Danish Patent office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Danish Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Danish Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Danish Patent Office or any person by the applicant in the individual case.

SWEDEN

The applicant hereby requests that, until the application has been laid open to public inspection (by the Swedish Patent Office), or has been finally decided upon by the Swedish Patent Office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the International Bureau before the expiration of 16 months from the priority date (preferably on the Form PCT/RO/134 reproduced in annex Z of Volume I of the PCT Applicant's Guide). If such a request has been filed by the applicant any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Swedish Patent Office or any person approved by a applicant in the individual case.

NETHERLANDS

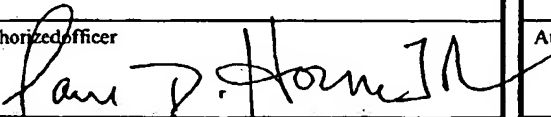
The applicant hereby requests that until the date of a grant of a Netherlands patent or until the date on which the application is refused or withdrawn or lapsed, the microorganism shall be made available as provided in the 31F(1) of the Patent Rules only by the issue of a sample to an expert. The request to this effect must be furnished by the applicant with the Netherlands Industrial Property Office before the date on which the application is made available to the public under Section 22C or Section 25 of the Patents Act of the Kingdom of the Netherlands, whichever of the two dates occurs earlier.

Applicant's or agent's file reference number	PA002PCT	International application No.	UNASSIGNED
---	----------	-------------------------------	------------

INDICATIONS RELATING TO A DEPOSITED MICROORGANISM

(PCT Rule 13bis)

A. The indications made below relate to the microorganism referred to in the description on page <u>79</u> , line <u>N/A</u>	
B. IDENTIFICATION OF DEPOSIT	
Further deposits are identified on an additional sheet <input type="checkbox"/>	
Name of depositary institution American Type Culture Collection	
Address of depositary institution (including postal code and country) 10801 University Boulevard Manassas, Virginia 20110-2209 United States of America	
Date of deposit 11 August 1999	Accession Number PTA-498
C. ADDITIONAL INDICATIONS (leave blank if not applicable)	
This information is continued on an additional sheet <input type="checkbox"/>	
D. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE (if the indications are not for all designated States)	
Europe In respect to those designations in which a European Patent is sought a sample of the deposited microorganism will be made available until the publication of the mention of the grant of the European patent or until the date on which application has been refused or withdrawn or is deemed to be withdrawn, only by the issue of such a sample to an expert nominated by the person requesting the sample (Rule 28 (4) EPC). Continued on the Attached Pages 2 & 3	
E. SEPARATE FURNISHING OF INDICATIONS (leave blank if not applicable)	
The indications listed below will be submitted to the International Bureau later (specify the general nature of the indications e.g., "Accession Number of Deposit")	

For receiving Office use only	For International Bureau use only
<input checked="" type="checkbox"/> This sheet was received with the international application	<input type="checkbox"/> This sheet was received by the International Bureau on:
Authorized officer 	Authorized officer

ATCC Deposit No. PTA-498**Page N . 2****CANADA**

The applicant requests that, until either a Canadian patent has been issued on the basis of an application or the application has been refused, or is abandoned and no longer subject to reinstatement, or is withdrawn, the Commissioner of Patents only authorizes the furnishing of a sample of the deposited biological material referred to in the application to an independent expert nominated by the Commissioner, the applicant must, by a written statement, inform the International Bureau accordingly before completion of technical preparations for publication of the international application.

NORWAY

The applicant hereby requests that the application has been laid open to public inspection (by the Norwegian Patent Office), or has been finally decided upon by the Norwegian Patent Office without having been laid open inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Norwegian Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Norwegian Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on the list of recognized experts drawn up by the Norwegian Patent Office or any person approved by the applicant in the individual case.

AUSTRALIA

The applicant hereby gives notice that the furnishing of a sample of a microorganism shall only be effected prior to the grant of a patent, or prior to the lapsing, refusal or withdrawal of the application, to a person who is a skilled addressee without an interest in the invention (Regulation 3.25(3) of the Australian Patents Regulations).

FINLAND

The applicant hereby requests that, until the application has been laid open to public inspection (by the National Board of Patents and Regulations), or has been finally decided upon by the National Board of Patents and Registration without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art.

UNITED KINGDOM

The applicant hereby requests that the furnishing of a sample of a microorganism shall only be made available to an expert. The request to this effect must be filed by the applicant with the International Bureau before the completion of the technical preparations for the international publication of the application.

ATCC Deposit No.: PTA-498**Page No. 3****DENMARK**

The applicant hereby requests that, until the application has been laid open to public inspection (by the Danish Patent Office), or has been finally decided upon by the Danish Patent office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the Danish Patent Office not later than at the time when the application is made available to the public under Sections 22 and 33(3) of the Danish Patents Act. If such a request has been filed by the applicant, any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Danish Patent Office or any person by the applicant in the individual case.

SWEDEN

The applicant hereby requests that, until the application has been laid open to public inspection (by the Swedish Patent Office), or has been finally decided upon by the Swedish Patent Office without having been laid open to public inspection, the furnishing of a sample shall only be effected to an expert in the art. The request to this effect shall be filed by the applicant with the International Bureau before the expiration of 16 months from the priority date (preferably on the Form PCT/RO/134 reproduced in annex Z of Volume I of the PCT Applicant's Guide). If such a request has been filed by the applicant any request made by a third party for the furnishing of a sample shall indicate the expert to be used. That expert may be any person entered on a list of recognized experts drawn up by the Swedish Patent Office or any person approved by a applicant in the individual case.

NETHERLANDS

The applicant hereby requests that until the date of a grant of a Netherlands patent or until the date on which the application is refused or withdrawn or lapsed, the microorganism shall be made available as provided in the 31F(1) of the Patent Rules only by the issue of a sample to an expert. The request to this effect must be furnished by the applicant with the Netherlands Industrial Property Office before the date on which the application is made available to the public under Section 22C or Section 25 of the Patents Act of the Kingdom of the Netherlands, whichever of the two dates occurs earlier.

What Is Claimed Is:

1. An isolated nucleic acid molecule comprising a polynucleotide having a nucleotide sequence at least 95% identical to a sequence selected from the group consisting of:

(a) a polynucleotide fragment of SEQ ID NO:X or a polynucleotide fragment of the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X;

(b) a polynucleotide encoding a polypeptide fragment of SEQ ID NO:Y or a polypeptide fragment encoded by the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X;

(c) a polynucleotide encoding a polypeptide domain of SEQ ID NO:Y or a polypeptide domain encoded by the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X;

(d) a polynucleotide encoding a polypeptide epitope of SEQ ID NO:Y or a polypeptide epitope encoded by the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X;

(e) a polynucleotide encoding a polypeptide of SEQ ID NO:Y or the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X, having biological activity;

(f) a polynucleotide which is a variant of SEQ ID NO:X;

(g) a polynucleotide which is an allelic variant of SEQ ID NO:X;

(h) a polynucleotide which encodes a species homologue of the SEQ ID NO:Y;

(i) a polynucleotide capable of hybridizing under stringent conditions to any one of the polynucleotides specified in (a)-(h), wherein said polynucleotide does not hybridize under stringent conditions to a nucleic acid molecule having a nucleotide sequence of only A residues or of only T residues.

2. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide fragment comprises a nucleotide sequence encoding a protein.

3. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide fragment comprises a nucleotide sequence encoding the sequence identified as SEQ ID NO:Y

or the polypeptide encoded by the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X.

4. The isolated nucleic acid molecule of claim 1, wherein the polynucleotide
5 fragment comprises the entire nucleotide sequence of SEQ ID NO:X or the cDNA sequence included in ATCC Deposit No.:Z, which is hybridizable to SEQ ID NO:X.

5. The isolated nucleic acid molecule of claim 2, wherein the nucleotide
10 sequence comprises sequential nucleotide deletions from either the C-terminus or the N-terminus.

6. The isolated nucleic acid molecule of claim 3, wherein the nucleotide
15 sequence comprises sequential nucleotide deletions from either the C-terminus or the N-terminus.

7. A recombinant vector comprising the isolated nucleic acid molecule of claim
1.

8. A method of making a recombinant host cell comprising the isolated nucleic
20 acid molecule of claim 1.

9. A recombinant host cell produced by the method of claim 8.

10. The recombinant host cell of claim 9 comprising vector sequences.

25 11. An isolated polypeptide comprising an amino acid sequence at least 95% identical to a sequence selected from the group consisting of:

(a) a polypeptide fragment of SEQ ID NO:Y or the encoded sequence included in ATCC Deposit No.:Z;

30 (b) a polypeptide fragment of SEQ ID NO:Y or the encoded sequence included in ATCC Deposit No.:Z, having biological activity;

(c) a polypeptide domain of SEQ ID NO:Y or the encoded sequence included in ATCC Deposit No.:Z;

(d) a polypeptide epitope of SEQ ID NO:Y or the encoded sequence included in ATCC Deposit No.:Z;

5 (e) a full length protein of SEQ ID NO:Y or the encoded sequence included in ATCC Deposit No.:Z;

(f) a variant of SEQ ID NO:Y;

(g) an allelic variant of SEQ ID NO:Y; or

(h) a species homologue of the SEQ ID NO:Y.

10

12. The isolated polypeptide of claim 11, wherein the full length protein comprises sequential amino acid deletions from either the C-terminus or the N-terminus.

13. An isolated antibody that binds specifically to the isolated polypeptide of
15 claim 11.

14. A recombinant host cell that expresses the isolated polypeptide of claim 11.

15. A method of making an isolated polypeptide comprising:

20 (a) culturing the recombinant host cell of claim 14 under conditions such that said polypeptide is expressed; and

(b) recovering said polypeptide.

16. The polypeptide produced by claim 15.

25

17. A method for preventing, treating, or ameliorating a medical condition, comprising administering to a mammalian subject a therapeutically effective amount of the polypeptide of claim 11 or the polynucleotide of claim 1.

30 18. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject comprising:

(a) determining the presence or absence of a mutation in the polynucleotide of claim 1; and

(b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or absence of said mutation.

5

19. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject comprising:

(a) determining the presence or amount of expression of the polypeptide of claim 11 in a biological sample; and

10

(b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or amount of expression of the polypeptide.

20. A method for identifying a binding partner to the polypeptide of claim 11 comprising:

15

(a) contacting the polypeptide of claim 11 with a binding partner; and

(b) determining whether the binding partner effects an activity of the polypeptide.

21. The gene corresponding to the cDNA sequence of SEQ ID NO:Y.

20

22. A method of identifying an activity in a biological assay, wherein the method comprises:

(a) expressing SEQ ID NO:X in a cell;

(b) isolating the supernatant;

(c) detecting an activity in a biological assay; and

25

(d) identifying the protein in the supernatant having the activity.

23. The product produced by the method of claim 20.

<110> Human Genome Sciences, Inc.

<120> 25 Prostate and Prostate Cancer Associated Proteins

<130> PA002PCT

<140> Unassigned

<141> 2000-07-19

<150> 60/144,972

<151> 1999-07-21

<150> 60/148,681

<151> 1999-08-13

<150> 60/149,173

<151> 1999-08-17

<150> 60/158,004

<151> 1999-10-06

<150> 60/194,689

<151> 2000-04-05

<160> 157

<170> PatentIn Ver. 2.0

<210> 1

<211> 733

<212> DNA

<213> Homo sapiens

<400> 1

gggatccgga	gccc aaatct	tctgacaaaa	ctcacacatg	cccaccgtgc	ccagcacctg	60
aattcgaggg	tgcaccgtca	gtcttcctct	tcccccaaa	acccaaggac	accctcatga	120
tctcccggac	tcctgaggtc	acatgcgtgg	tggtggacgt	aagccacgaa	gaccctgagg	180
tcaagttcaa	ctggtacgtg	gacggcgtgg	aggtgcataa	tgccaagaca	aagccgcggg	240
aggagcagta	caacagcacg	taccgtgtgg	tcagcgtcct	caccgtcctg	caccaggact	300
ggctgaatgg	caaggagtac	aagtgcagg	tctccaacaa	agccctccca	acccccatcg	360
agaaaaccat	ctccaaagcc	aaagggcagc	cccgagaacc	acaggtgtac	accctgcccc	420
catcccggga	tgagctgacc	aagaaccagg	tcagcctgac	ctgcctggtc	aaaggcttct	480
atccaaagca	catcgccgtg	gagtgggaga	gcaatgggca	gccggagaac	aactacaaga	540
ccacgcctcc	cgtgctggac	tccgacggct	ccttcttctc	ctacagcaag	ctcaccgtgg	600
acaagagcag	gtggcagcag	gggaacgtct	tctcatgctc	cgtgatgcat	gaggctctgc	660
acaaccacta	cacgcagaag	agcctctccc	tgtctccggg	taaatgagtg	cgacggccgc	720
gactctagag	gat					733

<210> 2

<211> 5

<212> PRT

<213> Homo sapiens

<220>

<221> Site

<222> (3)

<223> Xaa equals any of the twenty naturally occurring L-amino acids

<400> 2

Trp	Ser	Xaa	Trp	Ser
1				5

<210> 3

<211> 86

<212> DNA
 <213> Artificial Sequence
 <220>
 <221> Primer_Bind
 <223> Synthetic sequence with 4 tandem copies of the GAS binding site found in the IRF1 promoter (Rothman et al., Immunity 1:457-468 (1994)), 18 nucleotides complementary to the SV40 early promoter, and a Xho I site.

<400> 3
 gcgcctcgag atttccccga aatctagatt tccccgaaat gatttccccg aaatgatttc 60
 cccgaaatat ctgccatctc aattag 86

<210> 4
 <211> 27
 <212> DNA
 <213> Artificial Sequence
 <220>
 <221> Primer_Bind
 <223> Synthetic sequence complementary to the SV40 promoter and including a Hind III restriction site.

<400> 4
 gcggcaagct ttttgcaaag cctaggc 27

<210> 5
 <211> 271
 <212> DNA
 <213> Artificial Sequence
 <220>
 <221> Protein_Bind
 <223> Synthetic promoter for use in biological assays; includes GAS binding sites found in the IRF1 promoter (Rothman et al., Immunity 1:457-468 (1994)).

<400> 5
 ctcgagattt ccccgaaatc tagatttccc cgaaatgatt tccccgaaat gatttccccg 60
 aaatatctgc catctcaatt agtcagcaac catagtcccc cccctaactc cgcccatccc 120
 gccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat 180
 ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt 240
 ttttgagggc ctaggctttt gcaaaaagct t 271

<210> 6
 <211> 32
 <212> DNA
 <213> Artificial Sequence
 <220>
 <221> Primer_Bind
 <223> Synthetic primer complementary to human genomic EGR-1 promoter sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); including a Xho I restriction site.

<400> 6
 gcgctcgagg gatgacagcg atagaacccc gg 32

<210> 7
 <211> 31
 <212> DNA
 <213> Artificial Sequence
 <220>
 <221> Primer_Bind

<223> Synthetic primer complementary to human genomic EGR-1 promoter sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); including a Hind III restriction site.

<400> 7
gcgaagcttc gcgactcccc ggatccgcct c 31

<210> 8
<211> 12
<212> DNA
<213> Homo sapiens

<400> 8
ggggactttc cc 12

<210> 9
<211> 73
<212> DNA
<213> Artificial Sequence
<220>
<221> Primer_Bind
<223> Synthetic primer with 4 tandem copies of the NF-KB binding site (GGGGACTTTC), 18 nucleotides complementary to the 5' end of the SV40 early promoter sequence and a Xho I restriction site.

<400> 9
gcgccctcga ggggactttc ccggggactt tccggggact ttccgggact ttccatcctg 60
ccatctcaat tag 73

<210> 10
<211> 256
<212> DNA
<213> Artificial Sequence
<220>
<221> Protein_Bind
<223> Synthetic promoter for use in biological assays; includes NF-KB binding sites.

<400> 10
ctcgaggga ctttccggg gactttccgg ggactttccg ggactttcca tctgccatct 60
caattagtag gcaaccatag tcccgcctt aactccgcc atccgcctt taactccgc 120
cagttccgc catttccgc cccatggctg actaattttt tttatttatg cagaggccga 180
ggcgcctcg gcctctgagc tattccagaa gtagtgagga ggctttttt gaggcctagg 240
cttttgcaaa aagctt 256

<210> 11
<211> 1200
<212> DNA
<213> Homo sapiens

<400> 11
tcgacccacg cgccgcctt gtcataact cttagcatat ctttttttcc tttggctttg 60
catggctttt cttcaggtag tgtctcggtg tcattctgct aatcattgtt acagaatggt 120
gacttcattt gtgctaacag tacaacagca gatttggttc aggttaatc taagtgttaa 180
cttttttttc tgggtgcttt ttggattgat gactgtctca ctttgactat acccatgttt 240
tgcatgcaat gactcatgca tgggttttct aactagctaa tattaacaat ttattccata 300
taaaaatgga attttgcaac atcctttaat aagggtgagg aagcatgaac ctcagacttc 360
tggcactatt acatagtaag cacatgaagt agtttgataa taaatagcag ttctagtact 420
tcacatttca cccgtgtgtg caatgcctt ttctgggggg tggggggtga gggaaaacct 480
ggtagtgaat gtgtagtgtg ggaataaaga aaagcactaa atcctgcctt ttttgtgtgg 540
tttctttttg atacaactag gttattcata atgtatacct agaaaagtga aattgaaaat 600

acaaaaagat	gtatcatttt	tatttgaatc	catcatgcag	tgtacatttc	agataatttc	660
cttcagtcct	cagataggag	tgtatccaaa	catctaattt	tatgtgcact	gtgtatctta	720
tatgaatggt	ttattttata	taccacatgc	aaaaatgtcc	atatgcacta	tttaaagtgt	780
ttaaataata	tattccttct	ttataatgct	aaatctatat	gagtaccata	tttttataag	840
tcagtgggtc	gactggtttc	attttagaat	taacagctgc	ttcaatatgt	tattcaatgt	900
taatgttttg	ctgtgagtag	aatatgtaaa	agtggcatgg	cagcacttat	gctctgtgac	960
agtattgtgt	gtcatagttg	agcagtagct	ggtagaatta	ggcagttggt	gatagtttta	1020
ctttgggtaca	aataaaaaact	gtatatctat	atacaataaa	tatatagata	tatatgtcca	1080
ccagtataat	ggcattgctg	tgtctggcac	ttcattgtac	agacttttat	aataaaaagaa	1140
cttgaaagtt	ctaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	1200

<210> 12

<211> 1106

<212> DNA

<213> Homo sapiens

<400> 12

ggcacgaggt	ttccggttcc	cagagtgggg	cacagcgagg	cgctaggggg	aacgctggcc	60
tctgaaacta	gctctgggac	cggggtctgc	ggccggcccc	tagctggccc	cgtctcccat	120
ccccagaagg	gtattcactg	gggattctga	gctttggcta	ctccagtttc	ccacgacacg	180
atgttccctt	tctacagctg	ctggaggact	ggactgctac	tactactcct	ggctgtggca	240
gtgagagaat	cctggcgacac	agaagaaaaa	acttgcgact	tggtaggaga	aaagggtaaa	300
gagtcagaga	aagagttggc	tctagtgaag	aggctgaaac	cactgtttta	taaaagcttt	360
gagagcactg	tgggccaggg	ttcagacaca	tacatctaca	tcttcagggt	gtgccgggaa	420
gctggcaacc	acacttcttg	ggcaggcctg	gtgcaaatca	acaaaagtaa	tgggaaggag	480
acagtggtag	ggagactcaa	cgagactcac	atcttcaacg	gaagacaatt	ttaaccctgt	540
gtctgaggag	cgtggcaaa	tccaagattg	tttctacctc	tttgagatgg	atagcagcct	600
ggcctgttca	ccagagatct	cccacctcag	tgtgggttcc	atcttacttg	tcacgtgagt	660
atgccttcc	ttatcagaac	agaccttcc	tccttttgtt	tttaagggtca	gtaactatat	720
ggtggtacat	aagcacaat	tgtatgtgta	cattatgctg	taattgatgg	ggataacttt	780
ttaaatcctc	tggctataaa	ataattctgc	caggacacag	ggctcacgcc	tgtaatccca	840
gcacttttgg	aagccaaggc	agatggatca	cctaagggtca	ggagtttgag	accaacctga	900
ccaacatgat	gaaactccgt	ctctactaaa	aatacaaaat	tagccagcac	agtggcacat	960
gcctgtaatc	ccagctactt	gggaggctga	ggcagtagaa	tcacttgaac	ctgggaggtg	1020
gaggttgtag	tgagctgagg	tgcaccatt	gcactccagc	ctgggcaaca	agggcaaaac	1080
tccgtctaaa	aaaaaaaaaa	aaaaaa				1106

<210> 13

<211> 887

<212> DNA

<213> Homo sapiens

<400> 13

ccacgcgtcc	gagcagacac	aatggtaaga	atgggtgctg	tctgtctgtc	tctgtctgtg	60
cttctgggtc	ctgctgtccc	ccaggagaac	caagatggte	gttactctct	gacctatate	120
tacactgggc	tgtccaagca	tgttgaagac	gtccccgcgt	ttcaggccct	tggctcactc	180
aatgacctcc	agttctttag	atacaacagt	aaagacagga	agtctcagcc	catgggactc	240
tggagacagg	tggaaggaa	ggaggattgg	aagcaggaca	gccaacttca	gaaggccagg	300
gaggacatct	ttatggagac	cctgaaagac	atcgtggagt	attacaacga	cagttaacggg	360
tctcacgtat	tgcagggaag	gtttggttgt	gagatcgaga	ataacagaag	cagcggagca	420
ttctggaaat	attactatga	tggaaaggac	tacattgaat	tcaacaaaga	aatcccagcc	480
tgggtcccc	tgcacccagc	agccccctac	tcctgccacg	tgcagcacag	cagcctggcc	540
cagccccctg	tgggtgccctg	ggaggccagc	taggaagcaa	gggttggagg	caatgtggga	600
tctcagaccc	agtagctgcc	cttctgcct	gatgtgggag	ctgaaccaca	gaaatcacag	660
tcaatggatc	cacaaggcct	gaggagcagt	gtggggggac	agacaggagg	tggatttgga	720
gaccgaagac	tgggatgcct	gtcttgagta	gacttggacc	caaaaaatca	tctcaccttg	780
agccaccccc	cacccattg	tctaattctgt	agaagcta	aaataatcat	ccctccttgc	840
ctagcaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaa		887

<210> 14

<211> 1918

<212> DNA

<213> Homo sapiens

<400> 14

gggcccggact	gaggctctta	cagtgggtccc	tgctggccct	tggtgaccgg	tcgcctcagt	60
tccgaccg	accggtacgc	tgctgcgctg	acgtgggtcc	cggaagtagg	gctggcgtag	120
ggccgccatg	ttgcagcagg	atagtaatga	tgacactgaa	gatgtttcac	tgtttgatgc	180
ggaagaggag	acgactaata	gaccaagaaa	agccaaaatc	agacatccag	tagcatcggt	240
tttccactta	ttctttcgag	tcagtgcgaat	catcgtctat	cttctctgtg	agttgctcag	300
cagcagcttt	attacctgta	tggtgacaat	tatcttgttg	ttgtcgtgtg	acttttgggc	360
agtgaagaat	gtcacaggta	gactaatggt	tgccctacgt	tggtggaatc	acattgatga	420
agatggaaaag	agccattggg	tgtttgaatc	tagaaaggag	tcctctcaag	agaataaaac	480
tgtgtcagag	gctgaatcaa	gaatcttttg	gttgggactt	attgcctgtc	cagtactgtg	540
ggtgatattt	gccttttagtg	cactcttctc	cttcagagta	aagtgggttg	cggtgggtat	600
catgggtgtg	gtgctacaag	gtgccaacct	gtatggttac	atcaggtgta	aggtgcgcag	660
cagaaagcat	ttaaccagca	tggtactttc	atattttgga	aagcagtttt	taagacaaaa	720
cactggagat	cctgaataga	gaaagcttat	gtgctttgtt	acattgggga	780	
acaactgaag	agattcttga	ctcaaccttt	tagagcttag	tccatgttgc	aacgaggagt	840
gttggctttg	tttttccact	taaaaacttt	atttataaaa	aggaaaagta	gttttcatat	900
taagttttta	tttccctttc	agcagttggg	gctagaaagt	atgtgttggc	actagaaaca	960
ttgtcaagat	ttgttctgtg	gtgtaggtat	gcacattcca	taggtatgca	cacggccatg	1020
taatatcagt	atatcccaag	ttaatgaaag	tgttcattta	cataggtaat	ggagaccttt	1080
gcattttgat	ccatagaaca	taggaggatg	ttcttagtct	gtctcaaagc	tctatatgtt	1140
tacatattat	ttctgtagat	tgttttcagg	agaaagtttt	gcttctatgg	taagagttag	1200
cactttggct	tatgtataag	ttagaaataa	ttgttagttt	ttaatatgca	cttcgtgggg	1260
aaattttctta	gacgtatgca	agcaagtga	aacaattagg	gccagtggta	ttaactactt	1320
tataaaaattt	tatttttgtt	tgtaagaagt	catctactta	aggcccagtt	aatataagtg	1380
gaatcatcat	agtttaagga	ataccagag	attgctgctg	ttctatttat	tttacagaaa	1440
ggrtagctag	attgaaagct	cttcagtgga	ccttgagcta	atagatcttt	taccactaaa	1500
agagcattat	tctcatgtca	taatgagaat	aataatttac	atacttggca	taataaatgc	1560
ctaaaagaca	ttttattttc	tgaatctatt	tttttcttgc	tataatgggg	atattgtaaa	1620
ttatgcattt	gtattaatgg	tatttcttaa	agcaatctat	gtaactgtaa	attaaaccaa	1680
tctacaaact	attgtaggca	tctgtaaatt	ctgtttagg	tattataaac	tttgttgaag	1740
tcttaactcag	cagattatgt	tgtgaatata	tttgtacatt	gttaaaatag	tttttaagat	1800
tattttgttta	attgaataag	tgtcttattg	gagtgtatgc	tttgaagggtg	caaaacttta	1860
tatttgtata	aaattctact	atttacaagg	caaaaaaaaa	aaaaaaaaaa	aaaaaaaaa	1918

<210> 15

<211> 1129

<212> DNA

<213> Homo sapiens

<400> 15

tcgaccacg	cgctccgcagc	ttccgtccctc	tctgttctct	tacagcaccc	ccacctgccca	60
gagctgatcc	tccctaggcc	ctgcctaacc	ttgagttggc	ccccaatccc	tctggctgca	120
gaagtccctt	tacccccaat	gagaggagg	gcaggaccag	atctttttgag	agctgagggt	180
tgagggcatt	gagccaacac	acagatttgt	cgctctgtc	cccgaagaca	cctgcaccct	240
ccatgcggag	ccaagatggg	gaatggaact	gaggaagatt	ataactttgt	cttcaagggtg	300
gtgctgatcg	gcgaatcagg	tggtggggaag	accaatctac	tctcccgtat	cacgcgcaat	360
gagttcagcc	acgacagccg	caccaccatc	ggggttgagt	tctccacccg	cactgtgatg	420
ttgggcaccg	ctgctgtcaa	ggctcagatc	tgggacacag	ctggcctgga	gcggtaccga	480
gccatcacct	cggcgtaact	tcgtggtgca	gtgggggccc	tcctggtgtt	tgacctaaac	540
aagcaccaga	cctatgctgt	ggtggagcga	tggtctgaag	agctctatga	ccatgctgaa	600
gccacgatcg	tcgtcatgct	cgtgggtaac	aaaagtgacc	tcagccaggc	ccgggaagtg	660
cccactgagg	aggcccgaat	gttcgctgaa	aacaatggac	tgctcttctc	ggagacctca	720
gccttgagct	ctaccaatgt	tgagctagcc	tttgagactg	tcctgaaaga	aatcttttgcg	780
aagggtgtcca	agcagagaca	gaacagcatc	cggaccaatg	ccatcactct	gggcagtgcc	840
caggctggac	aggagcctgg	ccctgggggag	agagggcct	gttgcatcag	cctctgacct	900
tggccagcac	cacctgcccc	cactggcttt	ttggtgcccc	ttgtccccac	ttcagcccca	960
ggacctttcc	ttgccctttg	gttccagata	tcagactgtt	ccctgttcac	agcaccctca	1020
gggtcttaag	gtcttcatgc	cctatcacaa	atacctcttt	tatctgtcca	cccctcacag	1080
actaggaccc	tcaaataaag	ctgtttttata	tcaaaaaaaaa	aaaaaaaaaa		1129

<210> 16

<211> 969

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (2)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (5)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (15)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (37)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (118)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (254)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (255)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (839)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (887)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (897)

<223> n equals a,t,g, or c

<400> 16

gncantgacc	caacncaatt	aatgtgagtt	agctcantca	ttaggcaccc	ccaggcttta	60
cactttatgc	ttccggctcg	tatgttgtgt	ggaattgtga	gcggataaca	atttcacnca	120
ggaaacagct	atgaccatga	ttacgccaaag	ctcgaaatta	ccctcactaa	agggaaacaaa	180
agctggagct	ccaccgcggt	ggcggccgct	ctagaactag	tggatcccc	gggctgcagg	240
aattcggcac	gagnnctctc	acccctcggga	gacgctcgcc	cgacagcata	gtacttgccg	300
cccagccacg	cccgcgcgcc	agccaccatg	ctaggtaaca	agcgactggg	gctgtccgga	360
ctgaccctcg	ccctgtccct	gctcgtgtgc	ctgggtgcgc	tggccgagge	gtacccctcc	420
aagccggaca	acccgggcga	ggacgcacca	gcggaggaca	tggccagata	ctactcrgcg	480
ctgcgacact	acatcaacct	catcaccagg	cagagatatg	gaaaacgatc	yagcccagag	540
acactgattt	cagacctctt	gatgagagaa	agcacagaaa	atgttcccag	aactcggtt	600
gaagaccctg	caatgtgggtg	atgggaaatg	agacttgctc	tctggccttt	tcctattttc	660
agcccatatt	tcacgtgtga	aaacgagaaat	ccacccatcc	taccaatgca	tgcagccact	720
gtgctgaatt	ctgcaatggt	ttcctttgtc	atcattgtat	atatgtgtgt	ttaaataaag	780

tatcatgcat	tcaaaaaaaaa	aaaaaaaaaaa	ctcgrggggg	ggccccgtac	ccaattcgnc	840
ctatagttag	tcgtattaca	attcactggc	cgctcgttta	caacgtngtg	actgggnaaa	900
ccctggcggt	acccaactta	atcgcccttg	agcacatccc	cctttcgcca	gctggcgtaa	960
tagcgaaga						969

<210> 17

<211> 1098

<212> DNA

<213> Homo sapiens

<400> 17

ccacgcgtcc	gagagtccac	cttgcgaccg	tatccgctag	cgcgccctgg	gatgcgcttg	60
ggctccctgt	tcgttccac	atgcagggca	gcacaaggag	aatgggctgc	atgactgatg	120
tccacggcg	cttcctccag	ttgctgatga	cccatggcgt	gctagaggaa	tgggacgtga	180
agcgcttgca	gacgcactgc	tacaagggtcc	atgaccgcaa	tgccaccgta	gataagttgg	240
aggacttcat	caacaacatt	aacagtgtct	tggagtcctt	gtatattgag	ataaagagag	300
gagtcacgga	agatgatggg	agacccattt	atgcgttggg	gaatcttgct	acaacttcaa	360
tttccaaaat	ggctacggat	tttgagaga	atgaactgga	tttgtttaga	aaggctctgg	420
aactgattat	tgactcagaa	accggctttg	cgtcttccac	aaacatattg	aacctggttg	480
atcaacttaa	aggcaagaag	atgaggaaga	aggaagcgga	gcaggtgctg	cagaagtttg	540
ttcaaaacaa	gtggctgatt	gagaaggaag	gggagttcac	cctgcacggc	cgggccatcc	600
tggagatgga	acaatacatc	cgggagacgt	accccgacgc	ggtgaagatc	tgcaatatct	660
gtcacagcct	cctcatccag	ggtaaaagct	gcgaaacctg	tgggatcagg	atgcacttac	720
cctgcgtggc	caagtacttc	cagtcgaatg	ctgaaccgcy	ctgccccac	tgcaacgact	780
actggcccca	cgagatccca	aaagtcttcg	accctgagaa	ggagagggag	tctggtgtct	840
tgaaatcgaa	caaaaagtc	ctgcgggtcc	ggcagcatta	gccatcgctg	cctgctgagg	900
ggctggctgc	cttgagtggc	ctgatcgcca	cagcccttct	tggaaagaa	gcgtctgtgt	960
ttcaggttcc	acgcgagtca	cctctttcgt	cttaatgttc	accgtccaca	gctttggaat	1020
aaaccatcct	gggaagttaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	1080
aaaaaaaaaa	aaaaaagg					1098

<210> 18

<211> 1992

<212> DNA

<213> Homo sapiens

<400> 18

aattcggcac	gagctaagct	gcagtgatgt	tgcctatatt	taaattttct	caaattggcca	60
agctctgatg	gtctacttta	tttgagcaat	agttgagact	tataattgcc	tataaataaa	120
caaacaaatg	aactattttg	ttttttttct	cacaacatct	ggcctatatt	gtctgtcagg	180
aagccatggc	tcgaatgtaa	agtacatagt	tcttacatac	ttcaactgca	gctgggtccct	240
gacctcacca	ggtttcagag	atgttcttaa	aggaagccag	ctgtggcagg	tcacagattc	300
atgggaaatg	gaaagaacca	aggaatatag	ctcttgccct	acctttctac	ccactgcaga	360
tatagttcaa	gccagagtaa	tggagaact	taacttacta	gcctctcagg	ctgctcctat	420
ccctacctcc	cagtgtacag	cccccccca	tctcttttag	cccccttccc	tcacttcccc	480
ttttataatg	tcacacaaat	cagggacagt	aggatcacat	tataacctac	tttgtcatag	540
ggatttcgatt	tttcttatat	caaatcatgt	ttcctgaaac	ccagctgggg	catatgcact	600
caatgtctaa	tacatactta	ttaatgtacc	ggatattggc	cttgccccctg	gatatacagca	660
atatattata	aaaggttcca	gtagatgaga	cgattgagtc	tgaatacaat	tgcagtaaat	720
tgtgccataa	aagatattgt	actgttacgg	tcttagagtt	aaagccgctt	gaatgcagca	780
tgcacattca	tgtaaacaga	caatcagggg	aggcctagaa	taaccacaaa	aattctattg	840
gccttactgc	agccacctat	atgtagacaa	tggaggagat	agtttgtggg	ccattattgt	900
acctgttttc	atccatttag	atcagaatct	ctctttcagg	tcatttatta	aatatgattg	960
aaatgtttta	aagttcctga	acatgattca	tgatgattaa	aatatcatac	aactgataaa	1020
agactttaag	aactttatat	atttctctgt	gcctcaaaat	gtaacagaaa	ttattcttag	1080
agctttgatt	ttagctatcc	taattactgc	aaataaatat	ttgttcttat	agttttaaat	1140
caaaaagaaa	agtcttggtta	taaaacctta	agcttgaaat	catattaata	aaatatattg	1200
tacatagtgg	aaaattttca	gtagctaatt	taaaattttca	gaaaatgcta	ttaaagaatt	1260
ttgattcaag	tattttaaact	gtttagttat	gcatgcttct	tattaaccga	aaatgataat	1320
accatttagt	ttagtgtaca	gtatgagaag	caatacctaa	tctatgtgtg	ctattgtatt	1380
ttttcttagt	tgggtgtgct	gctcagaaaa	acataactgt	tatgtgtata	catacctgtg	1440
tatatataaa	aggtcaattt	atatattttt	ctataggaaa	atggagtaac	aagttcccta	1500
tctcccatat	ttatttgtcc	atagtaaaat	ggccacattg	atgataattt	ctagaactag	1560
tttctgagat	tgtcagccct	ttgtctaaaa	taatggcagt	attaatgatt	gacttctgtc	1620

actgccatag	ttacctggat	tgtcagcctt	ggtagccttt	gtctaaagtc	ctaaagagtt	1680
ccaaaaaaaa	tgtgttgaaa	tttaattgct	aaatagtggt	tgggtgattct	ttacagtagg	1740
aattgtaata	attttcttgc	aaataagtta	tttactgcta	ttgatattga	ataatttgct	1800
ttttattcag	atataattca	aaaagcatga	atataatgatt	attcataaat	tgtatacttt	1860
accagtaagt	tttcagagga	aataaagact	tttaaactct	aaaaaaaaaa	aaaaaaaaaa	1920
aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	1980
aaaaaaaaaa	aa					1992

<210> 19

<211> 1625

<212> DNA

<213> Homo sapiens

<400> 19

caagaacaaa	tctgaaggag	gcctctgaca	tcaagcttga	accaaatacg	ttgaatggct	60
ataaaagcag	tgtgacggaa	ccttgccccg	acagtgggtga	acagctgcag	ccagctcctg	120
tgctgcagga	ggaagaactg	gctcatgaga	ctgcacaaaa	aggggaggca	aagtgtcata	180
agagtgcac	aggcatgtcc	aaaaagaagt	cacgacaagg	aaaacttggt	aaacagtttg	240
caaaaataga	ggaatctact	ccagtgcacg	attctcctgg	aaaagacgac	gcggtaccag	300
atltgatggg	tccccattct	gaccaggggt	agcacagtgg	cactgtgggc	gtgcctgtga	360
gctacacaga	ctgtgctcct	tcaccgctcg	gttggttcagt	tgtgacatca	gatagcttca	420
aaacaaaaga	cagcttttaga	actgcaaaaa	gtaaaaagaa	gaggcgaatc	acaaggatatg	480
atgcacagtt	aatccttagaa	aataactctg	ggattcccaa	attgactctt	cgtaggcgtc	540
atgatagcag	cagcaaaaaca	aatgaccaag	agaatgatgg	aatgaactct	tccaaaataa	600
gcatcaagtt	aagcaaagac	catgacaacg	ataacaatct	ctatgtagca	aagcttaata	660
atggatttaa	ctcaggatca	ggcagtagtt	ctacaaaatt	aaaaatccag	ctaaaacgag	720
atgaggaaaa	taggggggtct	tatacagagg	ggcttcatga	aaatgggggtg	tgctgcagtg	780
atcctctttc	tctcttggag	tctcgaatgg	aggtggatga	ctatagtcag	tatgaggaag	840
aaagtacaga	tgattcctcc	tcttctgagg	gcgatgaaga	ggaggatgac	tatgatgatg	900
actttgaaga	cgattttatt	cctcttctct	cagctaagcg	cttgagggtta	atagttggaa	960
aagactctat	agataattgac	atttcttcaa	ggagaagaga	agatcagttc	ttaaggctta	1020
atgcctaagc	tcttgggtctt	aacttgacct	gggataacta	ctttaaagaa	ataaaaaatt	1080
ccagtcaatt	attcctcaac	tgaaagttta	gtggcagcac	ttctattgtc	ccttcactta	1140
tcagcatact	attgtagaaa	gtgtacagca	tactgactca	attcttaagt	ctgatttggtg	1200
caaatTTTTT	tcgtactttt	taaatagcct	tcttacgtgc	aattctgagt	tagaggtaaa	1260
gccctgttgt	aaaaaaaagg	ctcaagcaaa	attgtacagt	gatagcaact	ttccacacag	1320
gacgttgaaa	acagtaaatgt	ggctacacag	tttttttaac	tgtaagagca	tcagctggct	1380
ctttaatatata	tgactaaaca	ataatttaaa	acaaatcata	gtagcagcat	attaagggtt	1440
tctagtatgc	taatatcacc	agcaatgatc	tttggctttt	tgatttattt	gctagatggt	1500
tcccccttgg	agttttgtca	gtttcacact	gtttgctggc	ccagggtgtac	tgtttggtggc	1560
ctttgttaat	atcgcaaaacc	attggttggg	agtcagattg	gtttcttaaa	aaaaaaaaaa	1620
aaaaa						1625

<210> 20

<211> 2930

<212> DNA

<213> Homo sapiens

<400> 20

ggcacgagca	ccatctaagg	cagaggatgg	gatgtggtgg	ggcagggatc	tggtacagac	60
ataggcgctt	ccttccagga	acagtccctc	aggagaggca	gtggaccaga	agagatggcg	120
aaacgtgagc	tcgccggagg	tacagttttt	ccagggtgctt	aagccctccc	ccactgtgcc	180
acccttggtga	ccttcaggct	tgggtcatca	agatgcacag	catggaacac	aagctcctct	240
ggatcctgca	gctgggtgaca	tggaaactgtt	ttctgggtcca	catgaacaca	ggctccatcc	300
aggcccagct	cctacccacc	gcctctctgt	gggccagctg	ttcacagaag	gccttccatc	360
tgatgctccc	catgcctgc	ttgctctcca	gccgagtctg	gccaatttgc	catggggagg	420
ctgcagtgtc	caagcctgct	ggaaactggg	atgtagctgg	ggacgaaagg	acagacccaa	480
gcgttctccc	tgcttgagat	ggtgtggcca	cagcagtgga	aggctgcaca	caggcacatt	540
ccttcttcca	cagtggggca	ccaaggattc	tgctcctcatt	gctgggtaag	cagggagaag	600
agaagttttc	cccattgtcta	attttgggat	ttcagtggag	ccttttccat	ctgtccagga	660
gaacagaagg	gaaaaaaaga	tacttgaaag	aaactgaagg	aaatttaaac	aaagaaacac	720
ttgaaagaaa	ctggaaagaa	aaataatttt	tttatgtgaa	caaattttgc	aagaagaaaa	780
aagcataaaa	gacactaacg	gcaaactctat	gtttaaatgg	aaaatcgtct	aactggagaa	840
gggcggtatc	caccccatat	tcggatccca	gggtcctgag	gcctcgcatt	gagctggggg	900

tccctctga	gccccagtg	gtgtggaatc	agtgcactct	tgactgggccc	tgtagtaagg	960
tgctcatggg	gtttgtcttc	tcacccacca	tcagaggact	tttaaaatca	taggcgtaga	1020
gagttagcta	tctgctgaat	tactgccact	cttcttggtg	ggggctccta	gctgtggctg	1080
ggggctccag	gcgcccctgt	gattacctcc	tactgccacc	atggcgctca	ttcagattcc	1140
ccactctcac	taacattgct	tccttttttg	accagcagga	aacagcaggt	ctggccagat	1200
tctcacttgc	ccatcaatct	cgttcttgga	tgatttccct	cattgtgatg	cttctggggc	1260
acgttgacca	tatgcacctc	tagaacctaa	ccagggcttc	cttctaccag	ctgtggggcg	1320
gcttggctcg	gtaaccttgt	ctgctctgcc	attccactgc	tcctccatcc	actcgccaat	1380
cccaagagtc	tggcctccct	ccagccctgg	gcagactgac	cagcaagggtg	gacctttaca	1440
ttcaagcaca	gctggctttt	atgacataaa	gaactaaagg	ccgaaagaat	ctcttgctgc	1500
tgcaaagaac	agattttata	tttcttcctc	taatcttggc	aaatgacctt	taccttttgg	1560
aaagatttca	tattgcttcc	tcctccctgg	ataggacctc	atgtagcaca	gcgggactca	1620
aagaggagga	cattttctct	tgccagtgca	ctgggcagtg	gggctgtcct	tcaactgctg	1680
ctgccaaaat	tggttttcta	aaattcttcc	agtagagact	aaaagaagat	tcaattcctg	1740
taaccaaga	ctgagtctta	gggtccagt	ctccacctgc	ttggtttctt	atcctttgct	1800
gcctgcctgg	gggtggcctgg	aagcctgttc	agaaaggcac	aatgtggagc	ctgggtgtgc	1860
tccccacccc	caggaccgtc	aggtttacca	gtgtgtgcaa	tcgccatgta	ttcagaggga	1920
agtacctttg	ttacctacaa	cttaggagct	aggcctctgc	tacaagcact	tgaaaatgat	1980
atttttat	ttaacgtctc	aacaatctga	tatcggatgt	cgtttaacct	gggctcgtgg	2040
tagggctcca	gcatttctcc	ctccttctctg	gtttgcctgt	aggggtagac	tcggaagggtg	2100
ggtgggggtg	gcatttctctg	ttaggagtgt	atcagtgtct	gtcttattat	aagccctttt	2160
ctttgtgaa	tttgaagtag	caccaacaag	cctggattgt	gaaggattta	agaatcggtc	2220
tgtgggctac	tgagtgggtc	cttaggatac	tggcccagat	tttgccactg	ggtatggcag	2280
atcattttct	accatggcct	gctgctcttg	tagtggactt	cctgagtcca	atcccacctc	2340
ctggtgtaga	atttacactg	ctgcacctga	ggtcgatgtt	tcaaagtaag	atcaagccag	2400
tgttttgatc	tgggctctga	gcacaagtca	ggaaacacca	acatattcac	actctcccag	2460
taggttcctc	agtcggatgg	tgaatggcta	ttcgtaaatg	gctgggtctgg	ctctttgggtg	2520
ttggagcctt	tccaatagcc	ccatgaaaag	aagcatcacc	caaggatatt	gtaaaaagga	2580
tgtaacaagg	agatagggtg	gacattgtac	tcagtgggccc	ttggggccta	gcccagctct	2640
gagcagagga	ctgtggcatt	cactgtcctt	gagtgtttca	ccttcttgga	taacacacgg	2700
gccttctctt	ctggatttca	tcagagatta	cagccagatg	ggggctgaag	accatcctct	2760
tgaccacaga	ggtgtgactg	tgggaattcc	tcccaattta	tggttttcca	gaaaatctta	2820
gttcctttta	tttatagata	gcattgtctt	tgtgttaaga	aaccaaagag	aaataaagag	2880
aacactccta	ataaaaaaaaa	aaaaaaaaaaa	aaaaaaaaaaa	aaaaaaaaaaa		2930

<210> 21
 <211> 2655
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (22)
 <223> n equals a,t,g, or c

<400> 21						
ggattaaatc	caagtcgccc	cntattacgt	caatcscggt	atgcccgsct	gcattatgcc	60
agtcatgcct	tatggrrctt	ccacttgcag	acatctasgt	attagtcac	gtattaccat	120
ggtatcgggt	ttggcagtag	atcaatgggc	gtgatagcgg	tttgactcac	gggatttcca	180
agtctccacc	ccattgacgt	caatgggragt	ttgttttggc	acaaaaatca	cgggactttt	240
ccaaaatgts	gtamcaactc	cgccccattg	acgcaaattg	cggtaggcgt	gtacggtgga	300
aggctctatat	aagcagagct	cgtttagtga	accgtcagat	cgcttgagga	cgccatccac	360
gctgttttga	cctccataga	agacaccggg	accgatccag	cctccggact	ctagcctagg	420
ccgcgggacg	gataacaatt	tcacacagga	aacagctatg	accactaggc	ttttgcaaaa	480
agctattttag	gtgacactat	agaaggtagc	cctgcaggta	ccggctccgga	attcccgggt	540
cgacccacgc	gtccgatgca	gtttgcttgg	cagagctata	agcgttatgc	aatggggaaa	600
aacgaactcc	gtccactaac	aaaagatggc	tacgagggtg	acatgttcgg	aggcctcagc	660
ggggcaacag	tcattgactc	cctcgatacc	ctctacctca	tggagctgaa	ggaggagtct	720
caggaggcca	aggcctgggt	gggagagagc	ttccacctga	acgtgagcgg	agaagcatcc	780
tggtttgagg	tgaacatccg	ctacatcggg	ggactcctct	cagccttcta	cctgacagga	840
gaagagggtg	tccgaataaa	ggccatcagg	ctgggagaga	agctcctgcc	ggcggttcaac	900
acccccacgg	gaatcccaaa	gggcgtgggtg	agcttcaaaa	gtgggaactg	gggctggggc	960
acagccggca	gcagcagcat	cttggcggag	tttggatccc	tgcacttgga	attcttacac	1020
ctcactgaac	tctctggcaa	ccagggtcttc	gctgaaaagg	tcaggaacat	ccgcaagggtc	1080
ctcaggaaga	tcgaaaagcc	ctttggcctc	taccccaact	tcctcagccc	agtgagtggg	1140

aactgggtgc	aacaccatgt	ctcagttgga	ggactcgggg	acagttttta	tgaatatttg	1200
atcaaatect	ggttgatgtc	gggcaagaca	gatatggagg	ctaaaaatat	gtactacgaa	1260
gccttggagg	cgatagagac	ctacttgctg	aatgtctctc	ccgggggggt	gacctacatt	1320
gccgagtggc	gaggggggat	tctggaccac	aagatggggc	acctggcctg	tttctccggg	1380
ggcatgatcg	cccttggcgc	cgaggatgcc	aaggaagaaa	agagggccca	ctaccgagag	1440
ctcgcagccc	agatcaccaa	gacgtgtcac	gagtcatacg	cccgtcaga	caccaaactt	1500
gggcctgagg	ccttctgggt	taactccggc	agagaggccg	tggccaccca	gctgagcgag	1560
agctactaca	tctccggcc	agaggtgggt	gagagctaca	tgtacctgtg	gcgacagacc	1620
cacaacccca	tctacaggga	gtggggctgg	gaggtgggtg	tggccttggg	gaaatactgt	1680
cggacagaag	ccggtttctc	tgggatccaa	gacgtgtaca	gtagcacccc	caaccacgac	1740
aacaagcagc	agagcttctt	tctagcggag	acactaaagt	atctctatct	tctgttctct	1800
gaagatgact	tgctctccct	ggaagactgg	gtgttcaaca	ccgaggccca	cccactcccg	1860
gtgaaccact	cagacagctc	cggcagagcc	tggggcagac	actgacccca	tctcctgccg	1920
ccgccttggg	gccgcgcgag	ggatgccttg	ccttttcagg	atttgagact	gttctcaaaag	1980
ggattgggaa	cgaaggcccc	atctcgggca	gacccccagc	agatgtgtcg	gacaagcaac	2040
ttcttttctc	ctgtgaggag	acaagacttg	gagactcagc	gatgtcaggc	cagggccatg	2100
gccacactgg	cccacacatt	cctttctaca	gagaatttct	atgaagccca	ctcacttgcc	2160
attccagggc	caaaggaccg	gaggtttgca	tatccgcccc	ttgtatttga	tttgcttctc	2220
tttggtttct	tggtttttgt	ttttgcttga	ttttgtcttt	tctctacagt	ttagttttgt	2280
cacaattaca	catatagttt	tcaaaatcat	gcactttcta	aaatggtgtc	atcctgaaaa	2340
acaaaaccca	gtgtttgcac	acacacaaaa	tcttgacccc	gttatctata	ttttaaatgc	2400
tttttgccta	acactgaccc	tatgttcaac	tttgtgtcat	ttaccttata	atttgaggag	2460
gggtttccct	ttgggectca	gtgttacaaa	ttactagtgc	tattttcatt	attattgtaa	2520
tggaaaaatc	tgtggactag	aataaaagag	tttattgaat	aagaaaaaaa	aaaaaaaaaa	2580
aaaaaaaaaa	gggcggccgc	tctagaggat	ccctcgaggg	gcccacacta	cgcgtgcatg	2640
cgacgtcata	gctct					2655

<210> 22

<211> 1573

<212> DNA

<213> Homo sapiens

<400> 22

ccacgcgtcc	ggcggctggt	ttcttgggtga	gcccgggtcc	ctcaaggccg	gaaagaaagt	60
cgggcttctc	tagccccctg	aggactcgac	tcactgggtg	gcgatttagg	tccggaggag	120
gcgttgtgag	gtgagctttt	tcagaagcgc	gatcccagga	cacgtcggga	agcaagcatc	180
cccagagctg	cttggaaaga	ggaccaaaga	cgtctaaaaa	gtcatttggg	aatatctcta	240
aatatttggt	accatgtata	agctgctaaa	gagaaattgg	gcccacaaaa	actaattgaa	300
taattgaggc	agatttgtgt	gtatcatcaa	attctatcca	gaagttgaag	aatctgaatt	360
taaagattgt	gtgcatttaa	taagaggatg	acctttcagt	ttaatttcac	tatagaagac	420
catctggaaa	atgaattaac	acccattaga	gatggagctt	tgaccctgga	ttcctcaaaa	480
gagctgtcag	tctcagaaag	tcaaaaagga	gaagagaggg	acagaaaatg	ttctgcagaa	540
caatttgact	tgccctcagga	tcacttgtgg	gaacataagt	caatggaaaa	tgcagctccc	600
tctcaagaca	cagacagtc	actcagtcca	gccagcagtt	caaggaactt	ggagccacat	660
ggaaaacagc	cctccttgag	agctgccaaa	gagcatgcta	tgccataaga	tttaagaag	720
atgttagaaa	ataaagtcac	agaaacatta	ccaggtttcc	agcatgttaa	gttatcagta	780
gtgaaaacca	tcttgttgaa	agagaacttc	cctggagaaa	acatagtttc	aaaaagcttt	840
tcttctcact	ctgatctgat	tacagggtgt	tatgagggag	gcttaaaaaat	ctgggaatgt	900
acctttgacc	tcttggtcta	tttcacaaa	gccaaagtga	aatttgcctg	gaaaaaagtc	960
ttggatcttg	gttgtggatc	aggtttacta	ggtataactg	cattcaaggg	agggccaaa	1020
gaaattcact	ttcaagatta	taacagtatg	gtgattgatg	aagtaacctt	acctaatgta	1080
gtagctaact	ccactttgga	agatgaagaa	aatgatgtaa	atgagccaga	tgtgaaaaga	1140
tgcaggaaac	caaaagtaac	acaactatat	aaatgccgat	tttttcttgg	tgagtggctc	1200
gagttttgta	agcttgtact	aagtagtgaa	aaactttttg	taaaatatga	tctcattctc	1260
acctcagaaa	ccattttacaa	cccagattat	tatagtaatt	tgaccagac	tttccctaga	1320
ctgttaagta	aaaatggacg	tgtacttttg	gccagcaaa	cacattattt	tggtgtagg	1380
ggagggtgtc	atctctttca	gaagtttcta	gaagaaagag	atgtttttta	gaccagaata	1440
ctcaaaataa	ttgatgaagg	attgaagagg	ttcataattg	aaataacttt	taagtttctc	1500
ggttaattaa	cattcactga	gtatccaaaa	tgaataaac	agaaggacca	aaaaaaaaaa	1560
aaaaaaaaaa	aaa					1573

<210> 23

<211> 1998

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (1)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (2)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (33)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (90)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (101)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1710)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1852)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1873)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1947)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1953)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1989)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1991)

<223> n equals a,t,g, or c

<400> 23

nnaaccgggcc tttttacggg tcttgccctt ttngctgcct ttgggttcac atgttctttc
ctgcgttatc ccctgattct gtgaataccn gtattaccgc ntttgagtta agcttaatac
cgctcgccgc aagccgaacg accgagcgca agcgagtcag tgagcgagga agcggaagag

60
120
180

cgcccaatac	gcaaaccgcc	tctccccgcg	cggtggccga	ttcattaatg	cagctggcac	240
gacaggtttc	ccgactggaa	agcgggagct	gagcgcaacg	caattaatgt	gagttagctc	300
actcattagg	caccccaggc	tttacacttt	atgcttccgg	ctcgatgtt	gtgtggaatt	360
gtgagcggat	aacaatttca	cacaggaaac	agctatgacc	atgattacgc	caagctcgaa	420
attaaccctc	actaaaggga	acaaaagctg	gagctccacc	gcggtggcgg	ccgctctaga	480
actagtggat	cccccgkkct	gcagggaattc	ggcacgagcg	ggaatgggcg	tggcctgggc	540
ggggcgkcg	ctaggaccca	ccggagcgcc	gtgaacgtca	ccgagcgggc	ccgaggcccc	600
gggttgagcg	ggaggcgcg	tgggtccgg	cggtggctcc	ccgcggcggg	gccgggcccc	660
atctcgggcg	ggaaccgagc	gcagagccgg	tagcggaag	gatgaccacg	ctcacacgac	720
aagacctcaa	ctttggccaa	gtggtggccg	atgtgctctg	cgagttcctg	gaggtggctg	780
tgcattcat	cctctacgtg	cgcgaggtct	accccgtagg	catcttccag	aaacgcaaga	840
agtacaacgt	gccggtccag	atgtcctgcc	acccggagct	gaatcagtat	atccaggaca	900
cgctgcactg	cgtcaagcca	ctcctggaga	agaatgatgt	ggagaaagtg	gtggtggtga	960
ttttggataa	agagcaccgc	ccagtggaga	aattcgctct	tgagatcacc	cagcctccac	1020
tgtgtccat	cagctcagac	tcgtgttgt	ctcatgtgga	gcagctgctc	cgggccttca	1080
tctgaagat	cagcgtgtgc	gatgccgtcc	tggaccacaa	ccccccaggc	tgtaccttca	1140
cagtcctgg	gcacacgaga	gaagccgcca	ctcgcaacat	ggagaagatc	caggtcatca	1200
aggatttccc	ctggatcctg	gcggatgagc	aggatgtcca	catgcatgac	ccccggctga	1260
taccactaaa	aaccatgacg	tcggacattt	taaagatgca	gctttacgtg	gaagagcgcg	1320
ctcataaagg	cagctgaggg	ggcacctgcc	acccactga	tgcctaaact	gtcagacttt	1380
gggggatccc	cgcctagggc	agtgtgtcat	ggctgccctg	attccaagtg	ctcttatcgc	1440
ctctgtgtgt	ggatcgcccg	ccccagcccg	gggcccgtca	ggtctgcttg	gaggtatgct	1500
cccccaggcg	ggcagtgagg	gatgccgcaa	ctcgcacttc	tcagcctcct	gggggtccgc	1560
cgcccaacac	tgtctgtctc	aaatactgtg	ctgtgagttg	tttcaataaa	ggggccccaa	1620
gggctggaaa	aaaaaaaaaa	aaaaaaactc	gagggggggc	ccggtaccca	attcgcccta	1680
tagtgagtcg	tattacaatt	cactggccgn	cgttttacaa	cgctgtgact	gggaaaaccc	1740
tggcgttacc	caacttaate	gccttgacgc	acatccccct	tccgccagct	ggcgtaatat	1800
ccaaaaggcc	cgcaccgatac	gcccttccaa	cagttgcgca	acctgaatgg	cnaatggcaa	1860
attgtaaaagc	cgntaatatt	ttggtaaaaa	tcgcggtaaa	attttggtaa	atcaactcaa	1920
tttttaaac	aataggecca	aatcggnaaa	aancccttat	taaatcaaaa	gaattgacct	1980
agatagggnt	naatgggtg					1998

<210> 24

<211> 992

<212> DNA

<213> Homo sapiens

<400> 24

gccgtgggtg	ggtacaccaa	ctgcggaaag	accacgctga	tcaaggcact	gacgggagat	60
gccgccatcc	agccacggga	ccagctgttt	gccacgctgg	acgtcacggc	ccacgcgggc	120
acgtgccct	cacgatgac	cgtcctgtac	gtggacacca	tcggcttcc	ctcccagctg	180
ccgcacggcc	tcacgagtc	cttctccgcc	accctggaag	acgtggccca	ctcggatctc	240
atcttgacg	tgagggacgt	cagccacccc	gagggcgagc	tccagaaatg	cagcgttctg	300
tccacgtgc	gtggcctgca	gctgcccgcc	ccgctcctgg	actccatgg	ggagggtcac	360
aacaaggtgg	acctcgtgcc	cggttacagc	cccacggaac	cgaacgtcgt	gcccgtgtct	420
gccctgcggg	gccacgggct	ccaggagctg	aaagctgagc	tcgatgcggc	ggttttgaag	480
gcgacgggga	gacagatcct	cactctccgt	gtgaggctcg	caggggcgca	gctcagctgg	540
ctgtataagg	aggccacagt	tcaggaggtg	gacgtgatcc	ctgaggacgg	ggcggccgac	600
gtgagggtca	tcacagcaa	ctcagcctac	ggcaaattcc	ggaagctctt	tccaggatga	660
acggacgccc	acagaggcct	gcgggggtgg	ggcatcgctg	gctgggggagc	tgaggcggtta	720
ccgctgtgtt	gggggcagct	tggtgtcagg	tgcagcaggg	tcctccttgt	ctggttctgc	780
acccgtctcg	ctcccagcca	tttgcctggg	tgaccgtgca	ggccgggtgac	acggccgcac	840
ctgccccaaa	gcgggcgcgc	cgagcgtcca	ctccaagcct	gagcatccac	acaattccag	900
tgggcccctcg	gtgcctgctg	tgaactgctt	tcctcggaa	tgtttccgta	acaggacatt	960
aaacctttga	ttttaaaaaa	aaaaaaaaaa	aa			992

<210> 25

<211> 1926

<212> DNA

<213> Homo sapiens

<400> 25

tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	tttttttttt	60
tattttttatt	tttttttttt	taaaaatcca	ccaaaattac	ttaaatcatt	tgacagagtc	120

atacaattttt	aatgactat	aaaaatgtgc	agatcagtg	caaatttttt	ggcaaaggct	180
gaatggcaga	cagtaatcca	agacgcagac	aaccttgctt	ttaagccatc	cgttacacag	240
agaccagcag	gctcccttct	acctgtgaga	ccaaataatc	ccaaaatgta	accaaataat	300
cataactccc	cctactccta	aaacctaccc	ttcacccttc	actgtctacg	tcagcctctg	360
gacagacctg	atgcctgggg	tttggccatt	tccgccttcc	cacctgtgtg	tggctctcct	420
cctccctcct	caggctccct	agttctccca	ggcctaactt	ggcgtgcttc	cttctctctt	480
caccccaatc	aatccggctc	tccttccctt	tctgggttac	gccacctgcc	cgactcccgg	540
tttccccctc	ggggggccct	ccttgccctg	gggtctgctc	tccggccgcc	aagttctccc	600
aggggatccc	tccccacctg	cctccatata	gtcagactgt	gttcagtgtc	cgtttcgcgc	660
ccgtagcgac	accacgacag	aagcccacta	agtgcacctc	tgctcccaga	aacctcctg	720
gctcctccag	acctctcggg	gctctggccc	gcagctcccc	tctccctccg	gcggcggggg	780
tcggctgtga	ctcaccagag	ggccagcggc	cgggggtggg	gtgggggggc	cggaatgctc	840
tcacttctct	tcgcgcgacc	tgccccgagg	ctggagcgcc	gcccggccgg	gcctccggcg	900
gatgtttttc	ttgttcccaa	gcgtgtagtg	cgcgccagcc	gccccctccg	ggacctgcgc	960
gccagccacc	gcgccccag	gacccagcgg	gcctggctct	caccactcac	cccctcccca	1020
gcaggcacgc	atgcgggctc	gacccacacc	tctgcacctc	cccccaactt	ctgggagagg	1080
actcctggct	cagcccagcc	tctagctttc	cagaagcccc	tatatgccta	cctcatcttc	1140
gttataggcg	atgaaccttc	acttctgtcc	ccgtttccgc	acacgcacca	atccccactg	1200
gcatccctt	ccccttcggc	atcaccccca	ccttcatgcy	cccagctcc	ccactcacat	1260
ccgcccccta	ttgggctggc	tctagcttgc	aagagcagaa	ggtggcccag	ggcccagcct	1320
tccaggatgt	cccctggacc	ccctctctgg	gaaaggcgcc	agagctactg	gccactgaca	1380
agaccttttg	gcccctgtgc	tagacaggcc	tctgaatcca	cctgctccag	cccagagagc	1440
aggcccaggc	cgctgcctcc	acaccgcagc	cgacctcaga	gcacctgcc	tcaactgtga	1500
gttttttctt	ggcaatctct	tgaaacggaa	agggtatcag	tcctgactag	ggcgaacgtg	1560
aggctccatt	tgcatctcca	tccaccttag	ctgtctcctt	ggaaaaggac	agcaaggag	1620
tttgagggtg	actgacacag	tcatttctga	acagtgaaga	gatgacaggt	agggtgggtcc	1680
cccttgaaaa	gtcagccaga	agtagctgta	tctgatgcac	taggtggcag	ggagtgaagt	1740
ttggagtaaa	tgggactgaa	aacgtagaaa	aaccataata	atgccaaaga	gactatccct	1800
atggagaata	cctagaacat	gggtctttga	agtgtggaaa	aatgaaagaa	atccaatttg	1860
accagggcct	tagtgccaaa	gagatggcta	gcaggagagt	ggtgagaaag	cggacgcgtg	1920
ggtcga						1926

<210> 26

<211> 2542

<212> DNA

<213> Homo sapiens

<400> 26

ccacgcgtcc	ggtttatatt	tttttcacct	tttaaaca	aaattcttta	aaatatattta	60
atgcattctt	ttgaagaggt	agatgttttg	tacattttat	ggctcccaga	gcataatattc	120
agttggtgca	tggtgtggaa	gggggaattg	gaaattaaat	gaaaacctat	gactttgggtc	180
atgtcaatct	gtaagacaca	tcagtaaaag	ggtattatgc	tctgttggtt	ttgttttttt	240
gttttgcttt	tttttttttt	ttttcttttt	tggttttttg	tgatgtggct	taaatgcaat	300
agtttctttt	ttgggacata	tttctgccaa	ttaaagacta	gaagggcaca	actttttttt	360
taattaccat	agagaagata	cattaaaaaa	aatcttctga	tgttttgtag	ccataactaa	420
attatggtaa	aatgtgcac	tattgtgaaa	aggagcaacg	tagttttggg	ttttttgttg	480
tttggttggt	ttgctttggt	ttttaagaga	ttaaaatggt	tctggataag	gatttagcttc	540
tcgaagtgtc	catcattctg	tgtagaagct	taaatatgta	atgtaaccaa	actccagtat	600
taaaaatctc	tcatgttggt	ttctttatac	aaagcaagat	aacggcatat	aacactgcca	660
ttacatggca	aaatgtttgc	taccttagtt	taaaaaacaa	tctcaaacaa	aagacttgct	720
tcaagggtgt	tttaaatagc	agtgattcag	aatttttttt	aatgaaagta	taattgcact	780
aaccttcttc	ctgctgctct	gattctgcat	ttgtgggtact	tgtgactacg	ttttttcaaa	840
tatagataga	tttaagctgc	taattttttt	tttttttagta	atcactacta	tatcatgtct	900
tttactctgt	ttataatata	aagtattttt	ttaaagatat	agatatataa	ccttgtgctc	960
atgcaactta	gagtaacata	tacagacaaa	tgattgcatg	aggccatggt	tatatgtgtg	1020
actaataagg	cttgctcatga	ttaacataat	ccaggtagtg	catttctgaa	gagaatagtc	1080
atcaaattta	tatctcgaag	attttaatta	aggaattgct	tattgttgag	cttagcaaat	1140
taataacact	atttctgtca	ctaattat	tgaggccttt	tagtactaaa	attttaacct	1200
gtgttctaa	tagaaactga	tttaacccaa	gtaatgcagc	tttgattgat	ttcagcattc	1260
gttgctttgc	tattttttaca	aaacagcatt	gattgaagca	agtcttggtt	ttactaagggt	1320
agggtagacc	ttgtcattgg	taaagagaat	aaatacactt	aatttcacaa	tacattgtta	1380
tatgtacccc	agttgttggt	agtggggact	atgatactgt	aataaatatt	ttaaaaattt	1440
acatcaagag	aggcagtcac	tcacgatggg	tttgtgccag	ctcttttttag	ggttttggat	1500
cacattagag	atatttagaa	catattaccc	tgtgacttac	gtaggaaacc	taatatgctg	1560
agtatctggc	acttgaattc	ctgcttttat	tgctggagggt	ccacatctgt	ggttgacctc	1620

tggtattggt	taaaaaaaaat	aaataaaaaat	taaaaaaaatc	tgtgcaataa	ttttaaaatg	1680
tgctcccagg	aatagacaca	aatgttttga	gtatctttta	agctgcattt	tccttttagtg	1740
atgcatttgt	caattgcact	gaattttaa	ctgaaagtca	gaggtgatta	ttgatagtac	1800
ttttgtattt	tgatatggac	agtttattca	tttgcataca	gttattgact	ttttcccagc	1860
tgattaaaag	atagtcaaga	aattctgcaa	tatagctgcc	aaaatagaca	gctacatttt	1920
tatgatattg	tcattctttt	tgtttttttt	ttcttttttt	tccttagcta	ttttacttaa	1980
gcataatagc	cacaatagga	catataaaa	attataaata	cagagcttta	ttatcctgac	2040
gtcttgggtc	ttttaagtat	atacttttct	gaaaggatc	cattttgtag	gcttgggttc	2100
ttcatgagca	tacgattggt	tatttttgct	gctgttctca	acatcatcat	tgctgtctga	2160
tgtgccacga	tgctgctcca	atagacagca	ataagattgt	ctctaatttg	agcagtaaca	2220
tgattgcaag	agaccaagtt	tcacagcttg	taaagttctg	tatttgggat	tcttgcttat	2280
ttttccgcct	gtgtttttct	gagaacttat	tcctgatgat	caattgaatc	cagtagtttt	2340
tctatgctat	ttgttgttgt	ataagctact	gtaagaaact	tatcataagg	aaaaatagaa	2400
aggaaaaact	gaatcaatac	tcattgatta	aaatggaata	aagaaagagc	agctgccact	2460
tttaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	2520
aaaaaaaaaa	aaaaaaaaaa	aa				2542

<210> 27

<211> 1639

<212> DNA

<213> Homo sapiens

<400> 27

ggaacattac	aagggtttt	ataaaaaacc	ctttgttcat	atttcttccc	tttaaaatat	60
gtaatgtcaa	aaatgactca	ccttttaaaa	attatgcatg	aaaacagggtg	gtaaaccattc	120
agtaatacgc	tatttctcca	acatcaagac	aactaaaaca	aatgataaaa	atgttttattt	180
ttacactcca	gcataatcggg	tgagtttttag	ggatgtgtat	gaatatttaa	atcttttaaat	240
ttcagtttta	atgaaagctg	aacttaatag	gaaagctagc	tcttggtaac	tagcaatgat	300
caggcattgt	ttgcctctgt	cagggttttct	tatctgtttt	aggtacattt	tttcagattc	360
tgattgtttg	agttaatggt	tgaattttta	aagttttttag	ttacttaaaa	tatgatttta	420
aattacatata	taatttagaa	aattcctgtg	tttacttata	ttttaaattg	tgaaatggat	480
ccaatcatta	gaacagagag	aatagtctct	tgaactgaa	atactttagt	tttactgacc	540
ttgtgtaaag	ataatatgaa	gaaccagctt	ccaaaagaaa	ccagcatatg	gcactataaa	600
ctatttccatt	tgagcaccat	tctttaccat	ggatatatta	attatgtatt	atagtggagt	660
gatcatacag	ttcccccaaa	tgtgatgttt	cagtgtattt	atttaaaaca	taaagtagac	720
cagggtgcagt	agctcacgcc	tgtaatccca	gcactttggg	aagccaaggc	aggaggatta	780
ctgaggcca	ggagtttaag	accagccttg	gcaacatagc	gagaccccca	tctctacaaa	840
gaaatttttaa	aagattatcc	aggcatgggtg	gtgcatgcct	atgggtccag	ctactcgga	900
ggctgaagcg	gtagaatcac	ttgagcacag	gaggtcaagg	ctgcagggtga	gccatgatcg	960
tgtcactgca	ctccagccta	ggcaacagag	caagaccctg	tctcttaaaa	caaaaaacct	1020
cacacatacc	taaaataagt	atataatctgt	aaccaaaagga	acttaatttt	agagccaaat	1080
ttttctatct	tggtggattt	tttacagtgt	ttaagataga	cacttaagtg	ttgccgtagt	1140
atcaatattt	ctgtcattga	attttttctc	acccttatt	atttctcact	tacaatttct	1200
gagcatgatt	tttcataatt	cattatcttc	agatcattac	tataataaga	taccgtggct	1260
cttatgcaga	tattgcagca	agccttgttt	ggtttggatt	tttgcattcc	tttttctccc	1320
attaagggaa	ttaagttttg	ggatgatgga	ttaaaagcct	gtaaagaatt	gtttgggtct	1380
taaaagttgg	gctcatgggtg	gcaatatacct	tttaatgtgt	tttaacattt	tttaatcaac	1440
atcgtttaaa	aataatttac	taaggattag	attatttaat	gaatgtcttt	ttttctatat	1500
attgaattag	atgttatccc	aaaaaaaaata	tcttttctc	tcaaaatgaa	aatatatgtt	1560
caaagatgtt	ttatttttag	aatagttata	aaaagtatta	gtcttaaaaa	aaaaaaaaaa	1620
aaaaaaaaaa	aaaaaaaaaa					1639

<210> 28

<211> 1816

<212> DNA

<213> Homo sapiens

<400> 28

gtgtgtgtgt	aaattaccac	tgcaccgtga	acatgtgtac	attaccaggg	caagagctag	60
cctgggaaac	ctaagtctgc	acattttccg	ccgtgttgca	tgttttctgt	tctctgcctc	120
tgtgtgtgtg	caagacagag	agataggcta	tttgtcaagt	cagctagtgtg	cctaggtatc	180
tttgtctcac	atctggctgt	ttcctcctag	agaaccatcc	agttggcttt	ccaggcctgg	240
agggtgagcta	atggatgagt	gaatattagc	agtggtgtgt	cctcatctct	ttgaggattt	300
gcctcagagt	tcactaccaa	gggattttctg	gaactaggag	ccattcttta	catcagttct	360

tgaggcttct	ttgatatcag	gggcaaaatg	atcccttctc	ttttctttct	tatatcctgt	420
gctttgtctc	ctgggtgatt	tctcttcaag	tcagttgtgg	gaggtgccta	gaacaacgct	480
aacacggggc	tcagtaagtt	tggcagatat	agttgatttt	ggccaaggtt	gttgaacagt	540
ccagaaattt	cttgagagag	aaaagaaggt	agaatgtgtt	gttttctctt	tctgtatttc	600
ctcccccagt	cacctataga	ttcctggaat	tagagctagc	acttcacaga	acagagctgg	660
acttgctctg	aagcttacag	ctggaacctc	tgagtgtgta	tacgtgcaca	ctctagaatc	720
agatgagcca	gactgcattc	cagctcagca	cctttctcgc	tatgtgatct	tgtgcaagct	780
gcctgacctc	tgtaactatc	tcttttctca	actgtatgat	agggggaaat	ggtacccatt	840
tttaaaaagt	agctgtgaga	attaaatcca	ttattgaata	taaaagcagt	agagcctagt	900
gtataatgaa	cactcaataa	atgttacagt	tatttttcat	attgatggga	gtaccaataa	960
caagactggt	atcaatcgcg	taattcttta	cctccagctg	aactacaggg	tgataatggt	1020
tgaatggaga	gttttgtgtt	tggaaactca	gagtagaaga	cagatgacaa	ggatgctggg	1080
aacctccctg	accctctaata	atagtgtctc	gcaaaccagc	cagggaggat	aggataggtg	1140
ataaagcaat	taggaaatct	aatcagatcc	ctccaagtgg	caatgttttc	tgaagtctaa	1200
ggacttaaga	attgaactaa	tttctctaaa	acaccattca	tggctctcca	ttcctgatat	1260
caagatttta	tgagtcttgc	atttctcatt	ctcatttgag	tgcactctgg	aattatggat	1320
tttatatttg	ctggcggttt	accataaata	agccttgggt	cttgtgaatt	catgtgtttt	1380
gaaaaacatc	gcttttgttt	agctttccag	tctgggtaca	ataatctctt	ctttctagaa	1440
gtcattccag	ggcaagtctt	caggatattt	ggtttctatg	ataaggtgat	aaacatggat	1500
ggcttgatct	tggagccacc	aaactacatt	gagctttaa	ttctccattg	ggtacaatgg	1560
ttacatgctt	cttaccctaa	aactctggag	ttacaagtca	tagttgtatg	taagtattaa	1620
attcactgaa	tggagctttt	ctctagacag	atctccttct	tgaatctggg	tttctgtgct	1680
gggtgcaatg	gctcactcct	gtacttacaa	caatttgtaa	ggcccaggca	gaaggattgc	1740
ttgaagccaa	gagttcgaga	ccagcctggg	caggaaagtg	agacctcatc	tctaatttaa	1800
aaaaaaaaaa	aaaaaa					1816

<210> 29

<211> 3147

<212> DNA

<213> Homo sapiens.

<400> 29

tttagagaga	tggtgtcttc	cagcaatctg	ccacaagggg	ggttagaggt	ccaggggata	60
ccggaagggg	gggatgggtg	agcaggatgg	tatcttccag	gaataaacc	tgccaggact	120
gctaggcggt	ttgcttatct	ttttgtgaat	atcaatgtga	cctctgagcc	tcacgaagtt	180
cttgccctgt	ggttcttgtg	gtatgtgaag	cagtgcgggg	gcaccactcg	gatattctct	240
gttaccatag	gtggccagga	acggaagttt	gtaggtggat	ctgggtcaagt	gagcgaacgg	300
ataatggacc	tcctcggaga	ccaagtgaag	ctgaaccatc	ctgtcactca	cgttgaccag	360
tcaagtgaca	acatcatcat	agagacgctg	aaccatgaac	attatgagtg	caaatacgta	420
attaatgcga	tcctccgac	cttgactgcc	aagattcact	tcagaccaga	gcttccagca	480
gagagaaacc	agttaattca	gcgtcttcca	atgggagctg	tcattaagtg	catgatgtat	540
tacaaggagg	ccttctggaa	gaagaaggat	tactgtggct	gcatgatcat	tgaagtga	600
gatgtcccaa	tttcaataac	cttggatgac	accaagccag	atgggtcact	gcctgccatc	660
atgggcttca	ttcttgcccg	gaaagctgat	cgacttgcta	agctacataa	ggaaataagg	720
aagaagaaaa	tctgtgagct	ctatgccaaa	gtgctgggat	cccaagaagc	tttacatcca	780
gtgcattatg	aagagaagaa	ctgggtgtgag	gagcagtact	ctgggggctg	ctacacggcc	840
tacttccctc	ctgggatcat	gactcaatat	ggaagggtga	ttcgtcaacc	cgtagggcagg	900
attttctttg	cgggcacaga	gactgccaca	aagtggagcg	gctacatgga	aggggcagtt	960
gaggctggag	aacgagcagc	tagggaggtc	ttaaatggct	tcgggaaggt	gaccgagaaa	1020
gacatctggg	tacaagaacc	tgaatcaaag	gacgttccag	cggtagaaat	caccacacc	1080
ttctgggaaa	ggaacctgcc	ctctgtttct	ggcctgctga	agatcatagg	attttccaca	1140
tcagtaactg	ccctgggggt	tgtgctgtac	aaatacaagc	tcctgccacg	gtcttgaagt	1200
tctgttttta	tgtctctctg	tcaactggtt	tcaataccac	caagaggaaa	atattgacaa	1260
gtttaaaggg	tgtgtcattg	ggccatgttt	aagtgtactg	gatttaacta	cctttggctt	1320
aattccaatc	attgtttaaag	taaaaacaa	tcaaagaatc	acctaattaa	tttcagtaag	1380
atcaagctcc	atcttatttg	tcagtgtaga	tcaactcatg	ttaattgata	gaataaagcc	1440
ttgtgatcac	tttctgaaat	tcacaaaagt	aaacgtgatg	tgtcatcag	aaacaatttc	1500
tgtgtcctgt	ttttattccc	ttcaatgcaa	aatacatgat	gatttcagaa	acaaagcatt	1560
tgactttctg	tctgtggagg	tggagttagt	gaaggcccag	cctgtaactg	tcctttttct	1620
tccttaggc	aatgggtgaac	tgtcattaca	gagcctagag	gctcacagcc	tcctggagga	1680
agcagcctcc	actttggatc	aggaaatagt	aaaggaaaag	agtgttgggg	gtagcggcat	1740
gcagaccctc	agaccagaat	ggggacatct	tgtgggtctg	tgccctcagg	atctcctgac	1800
cacttgtagt	ccctccgact	tctctagaca	tctagtctca	gtgctagctt	atttgtattt	1860
ttcctctttc	acttcttatg	gaggagagtg	tttaactgag	ttagaatgtt	gaaactgact	1920
tgtgtgact	tatgtgcagc	tttccagtgt	agcagaggaa	aatagtggca	ggactgtccc	1980

ccaggaggac	tccctgctta	gctctgtggg	agaccaacta	cgactggcat	cttctcttcc	2040
ccctggaagg	cagctagaca	ccaatggatc	cttgtcagtt	gtaacattct	atttcaactt	2100
caggaagca	gcagttttct	tttaattttt	cctatgacca	taaaattaga	catacctctc	2160
aacttacata	tgtcttcaac	atggttacct	ctgcataaat	attagcaaag	catgccaatt	2220
tctcttaagt	actgaaatac	atatgataaa	tttgactggt	atttggtgag	actatcagac	2280
agaaaagaaa	ttagggctct	aatttcctta	aagcaagctc	acttgcttta	gttgtaagt	2340
tttataaaaag	acatgaaatt	gagtcatttt	atatatgaaa	actaagttct	ctatcttagg	2400
agtaatgtcg	gcccacaagg	gtgcccacct	cttgttttcc	ccttttaaaa	actcagattt	2460
ttaaaagccc	tttccaaagg	tttcaactgt	aaaatacttc	tttttacaat	gtatcaacat	2520
attttttattt	aaggggaatt	aacaattgcc	agggaaacca	gccaacccaa	gtttattata	2580
tcattaacct	tatcataaat	tcaaacctaa	gttgctggac	cctggtgtga	ggacataaat	2640
cttccaaagt	tttgccctatc	ctaagagctg	catttttcta	ctgctcttta	ccttgcaattt	2700
tagctaattt	aggagttttg	agaatgtatt	ggatacgtc	cagtacataa	ggagttgccg	2760
catattatat	cagactgctt	tgagaaatct	cateccctagt	ctattgcagt	tgtttctatt	2820
agcttactga	ttaactcagt	cctgacacac	cttttgggaa	atgctgattt	aaacttctta	2880
actggcaaca	gttggaacag	taatcagttt	gctaacatat	ttaaagtctt	gaatgttgaa	2940
gaactcatgt	gatttaccct	tttcaacttt	ttggaaaacg	atttaattta	atccaattag	3000
attaacccta	ttaaactctg	ggttgggtat	ccaaatgaat	gccagtcgga	tgttgccaga	3060
cacgaaattg	ggagccaggg	atctcacgaa	atgcagttca	tcccacgcgg	aggtagcaca	3120
agccttttgc	tcttagccga	gagatga				3147

<210> 30

<211> 1109

<212> DNA

<213> Homo sapiens

<400> 30

tcgaccacg	cgctccgaaac	tttcttttct	ccccctttt	tttgcccaca	aatggtatta	60
taatgcttgc	ttagtcaaag	aagagagact	aaacaaggg	aaaaatttta	acagtacaga	120
atttgccatc	atatcattgc	cttgattcta	actgtttgtg	tcctaagatg	caaaagaagt	180
cagtggcttt	taactgttta	caaatagaa	gtgattgtaa	aatgtacagt	ttggttgtgt	240
ttgaattatg	aaatttcttc	agatataata	aaccatgact	ttttggctgc	tcaacattaa	300
ttgtctctct	tttgtgaatt	tatttgtagg	ctctttttta	taatgaaagt	ttcaaagttg	360
ctatgtatga	gggttctcat	agagcaaccg	attaaaaatc	taagcaaata	tttgaacatt	420
ttatctgaac	tcatcacaat	ttcaccctga	aataatgtga	gaacaatggg	aaactgtagc	480
ttgctctctc	ccaccctctc	tgagcatctt	tgggatcttg	ttgctcaaaa	ctcttctgtg	540
acttcatctt	ccccaccatt	tgtgcccate	tcaagcctca	gcaagaaacc	atgtggaaca	600
tgaagcttaa	tgacttgaca	gtgtactagt	gttaaacctc	catacctctg	ttacaaagcg	660
agaaacgcca	caccgggact	ggccttttct	tcccccttca	cggccctcgc	ttctccctgc	720
aggagctcgg	gggcgaaacc	tgtgtatgga	tttcagtgtg	tgacttcaga	tcatgtccca	780
acttgccagg	tgtgagctaa	tgttgtcgga	caccttacta	taagcaaagt	ttattcagtg	840
cgttcaatgt	atattgactt	ccatactggt	ttttccaaaa	accaaaggta	gctttgaaaa	900
accatgtctg	gaaatgtttg	gagcgttaag	ctgattgacc	ttctgacctt	ggggctttga	960
tagtataata	attcataact	gcgttaattg	tattgtttaa	gtgtttggga	gttttttgcg	1020
cttgttatgt	ggaaataaag	tgtttgattt	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	1080
aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa				1109

<210> 31

<211> 2324

<212> DNA

<213> Homo sapiens

<400> 31

ggcacgagtg	ggcaggtaga	gcccagctc	caggtgaggc	ctggccctgg	gcagggctctg	60
tggctgcgcc	cctcaggcca	gcagttgagg	tccatcgggc	tggccccagc	ccatctgctg	120
gcacagtagc	ctgggtgttg	gacagcagga	taggggtcta	aaggtgggtt	tctatccaaa	180
cgacaaaaaa	accaacagta	acaccagtga	aacccacac	tgctgggctt	ataaaaaatc	240
gtgccatcat	ggtgatttta	tccaagactg	ctccacttac	cccagtgctg	gggacaagtt	300
tctgttgaaa	ctttagatag	cagaattatt	tgcaatttgt	agcatagaaa	agatttttaa	360
atttttttac	aaaagggttt	taaacagatt	agggtaggtg	atggttttaa	tcaattaaagt	420
ggcattggaa	acctagggtt	tccttttgat	taagagcctt	ttttgtttct	gctctttgtc	480
agctttcagg	ggagaaggag	gccactggaa	aattatttcc	ctaagtgacg	gctgttgact	540
gcgtatgcca	aaaagggaca	ggaggcatgg	gatagcaggt	ctggtgacac	agctagggtc	600
ttcctagcag	ctcctcctcc	tccctcccaa	ggccccagg	aatcccttcc	tcccatgtcc	660

tggcagcagg	accccaggct	acatatggaa	ggtagagatg	tgggggtcct	gtgtcctgga	720
gtattatgtc	tccccacctt	ctgcagtttt	ctctgaacat	gtatgttgcc	catggtggga	780
gcgtgggtcac	tgtgcagttg	tgcacagatg	tctttccttt	accgttggcc	tttctgtctg	840
cctctccttc	ctctctgcag	cccaaatgga	aaacaattat	ttactccatt	ggaggggaaag	900
gaagagtctt	agaattccta	agggaaacctt	agcataaagg	ttttggggaa	ggaggccgta	960
ggccggcccg	gaggaagcaa	ttccacttgg	tttgacaact	tctgccactc	ccatgtcaga	1020
tgacttgcac	ttcttaaaga	gattgtctta	taacactaag	acatcctttc	taaagattca	1080
agtggacttg	actaagctga	gggtccacga	aatagaatat	gacatgtgag	ctgttttttg	1140
aaaacgaaga	tggagagagc	acttccccgt	aacgaaagca	aagtggtaag	cacaggggtga	1200
gaccctttta	cacagaatgg	tggagagaaa	agagaatgct	gaaaagtggc	tcagatgcag	1260
agtgttctgt	ggagaaactg	cagccccact	tctgtttccc	tggagtctcc	caatggatca	1320
ttcaggagtg	tcctatgtga	gaattgagcc	aaggaaaata	ctcatgcaac	cagcctgagt	1380
cgcgggtgagg	ggacgagagg	ttgtacacac	attggtagtt	atthttgcacc	agcagtgcct	1440
ttctcactgg	gggtacttgg	accctcagat	cttcttttct	aatagccatt	tgccacccca	1500
agtggtagtg	cggccatttc	tccttaaaac	accttcctta	cctttcccat	gtactcagtt	1560
tagctctcaa	agaaggggtg	aatcataaag	ccagtgaana	tttcaccctc	tgaggggagt	1620
ccccaatctg	aaggggaaga	gggtgacctc	agcggctttt	ctcccaaaaa	tcggctgaag	1680
gctgggtgtg	gatccttggt	cctctcctga	ccccatctgg	ctgctgcccc	gtctcccacc	1740
cctgtccccg	gggtctcgctg	gccctgcact	ccgccttagt	cctggggccg	gcgacacagt	1800
gggggctcct	cacttgctgc	agtgtcatag	caataaaatg	tgattcttgg	ggtcccccca	1860
gggagctgcc	catggcttta	tttatgaacc	tggttttcgg	gagtcagggg	aggagatgac	1920
tttgcttctg	tgcacagccc	cgtcttccag	gagccacgac	tcagaagaaa	aggggtgctca	1980
gacttttgtt	atacacatth	gctttgtgta	aataaaatgt	tacaatttta	tatgaaagat	2040
ggaataagcg	ctagagcttc	caactgtata	ttttttactt	ttatagattt	taaaactatg	2100
atcctttata	tgtgtgtttt	gggggagcta	tgataagttt	tatggcaaac	ggttggtatt	2160
gttaactttt	tattgtcatc	aaaagttcat	aaaagtccta	ttaatcccca	tattcttcta	2220
ctgcccttaa	ctctgggtata	caccaaaaag	aaatcctttac	tttccttggt	ttatcattat	2280
aaaaataaag	tatttttgcta	gtatggaaaa	aaaaaaaaaa	aaaa		2324

<210> 32

<211> 1255

<212> DNA

<213> Homo sapiens

<400> 32

ggtgaaatga	gacaccctac	cttcaattgt	tcatacagtg	gtaaaacaaa	ttctgatgta	60
cattcaggac	aaatgattag	ccctaaatga	aactgtaata	atttcagtg	aaactcaatc	120
tgthttttacc	tttaaacagt	gaatttttaca	tgaatgaatg	ggttctttcac	ttttttttta	180
gtatgagaaa	attatacagt	gcttaatttt	cagagattct	ttccatatgt	tactaaaaaa	240
tgthtttggtc	agcctaacat	actgagtttt	ttttaacttt	ctaaattatt	gaatttccat	300
catgcattca	tccaaaatta	aggcagactg	tttggtattc	tccagtggcc	agatgagcta	360
aattaaatca	caaaagcaga	tgcttttgta	tgatctccaa	attgccaact	ttaaggaaat	420
attctcttga	aattgtcttt	aaagatcttt	tgcagctttg	cagataccca	gactgagctg	480
gaactggaat	ttgtcttctt	attgactcta	cttcttttaa	agcggctgcc	cattacattc	540
ctcagctgtc	cttgcagtta	gggtgtacatg	tgactgagtg	ttggccagtg	agatgaagtc	600
tcctcaaaag	aaggcagcat	gtgtcctttt	tcataccttc	atcttgctgc	tgggattgtg	660
gatataacag	gagccctggc	agctgtctcc	agaggatcaa	agccacaccc	aaagagtaag	720
gcagattaga	gaccagaaag	accttgacta	cttccctact	tccactgctt	tttctgcata	780
tttaagccatt	gtaaatctgg	gtgtgtttaca	tgaagtgaana	attaattctt	tctgcccttc	840
agttctttat	cctgatacca	tttaacactg	tctgaattaa	ctagactgca	ataattcttt	900
cttttgaaag	cttttaaaag	ataatgtgca	attcacatta	aaattgattt	tccattgtca	960
attagttata	ctcattttcc	tgccttgatc	tttcattaga	tattttgtat	ctgcttggaa	1020
tatatattct	tcttttttaac	tgtgttaattg	gtaattacta	aaactctgta	atctccaaaa	1080
tattgtcatc	aaattacaca	ccatgttttc	tatcattctc	atagatctgc	cttataaaca	1140
tttaaatata	aagtactatt	taatgattta	acttctgttt	tgaatgtgtg	tatacacgtg	1200
gatttttttc	tcattaaata	ataattctag	tatttgaaaa	aaaaaaaaaa	aaaaa	1255

<210> 33

<211> 534

<212> DNA

<213> Homo sapiens

<400> 33

ggcacgagca	gaagaggaag	tcagagcgat	gtgctgtgaa	atctactacc	gtttgctggt	60
------------	------------	------------	------------	------------	------------	----

tttgaaaatg	gagaaaaaga	gtgaggaact	gagaaacatg	gatggccttg	ggaacgtgga	120
aaagggtcac	tgaaatggga	cgacatgaac	tcaaggaggc	tatttatgac	catgtcattt	180
gcaacatgaa	gaaagcttat	ctggagtga	agtaaatgag	accaacagag	ataagagacc	240
cggagaaatc	ctggttacac	tgcttgaatc	ctgtcagtc	tatactggag	tcctgttaat	300
acaaaataat	agtaataatc	cctctgtttc	ttatgtttat	gccaacttca	acaaaaagaa	360
acttgactaa	gagacaatat	aagaacttaa	tgtgtaatta	agaaagaact	ctccaccacg	420
gggaatgtga	aaggtatatg	agtcctttt	cacgatgcga	tgcatgtct	tttaaataag	480
ccatacttta	tgttcaataa	aaagagaata	agcaggaaaa	aaaaaaaaaa	aaaa	534

<210> 34

<211> 1756

<212> DNA

<213> Homo sapiens

<400> 34

ttgacttctt	tctatagagg	gccacttcta	aagctacagc	attcatttgg	aaaaggacac	60
tgggatcaac	acgtaagcgt	tgcaagcaca	ggggccgcct	ctcttgacaga	caggtggcca	120
aagcagggtc	tgtgctgggc	cagaagtgga	agtaattcct	cgccagctac	acattcagtc	180
tgactgggtg	atgattggga	gtgtttgtcc	ctccctcccc	caataattga	tggccttgag	240
atctgccagc	atctcaaagg	cagattcgtg	gctctgttcc	cagacttagg	tctcagttat	300
ttaatgtgta	aatgacacaa	tcaagagact	caacactaat	tggaatgctg	atttcaagta	360
tttctctggc	tagctgggtg	aagccataga	ttgtggaatg	tgatttccat	ttctgaatca	420
gtgggtggca	ggttgggttac	agaaatgctt	cctattagtc	tggaggccct	ctaggagtct	480
gacatgtaat	taacaaagca	aaaggtaaac	atgaagggtc	ggcatggagc	tgggtgtata	540
tctggctgtg	ggacgtcggt	ttcttttctc	cactgcaaat	gagactacag	ttgtgatgat	600
ggtgggaggt	tccaggggtg	caggtcttca	ccaaggacca	tggtgtacca	cagcctctgc	660
tgagctgagg	gacacatgtc	cccctgcttt	cccaacacca	ccaactttgg	atttcacaga	720
ttgtctctag	gctgggggtc	gccttgactg	tattacacct	ggctggctcc	tttaatcaca	780
gcattctaga	aatccatatt	ggatgtggaa	taccagaatc	ctatgtagca	tcctagtttg	840
caatccaatt	atagagttct	ttaaaaattc	tggagtctct	tatgagaact	agaaccatga	900
tgaggtcaga	ggcaggcact	ttgcccacag	ttgatcctgg	aggcctccag	actctgtgag	960
gtgagcacct	ggcagcacct	ctgttctctc	tttgaggcca	gtaattgcct	ggggaagaga	1020
gacagggaaa	tggaaagcat	gatcagggca	aatcagccag	catttctagt	ttggagggtc	1080
acctggcctt	ttccgtgggc	tcagggccat	ctgaaacact	gccctgtaaa	gctggtgctg	1140
ggctgtcctt	gtgcctggag	agtgtctaaa	ttgacatttc	aaatacccag	agagcagggg	1200
gagatacccc	gcattgtccat	agctgcaaag	aaatgtctag	gggggtctcc	tcttctgact	1260
cctcacctgg	cgatgtatca	gcattcaata	ctgaacactc	ttaggggccc	aagtatggcc	1320
tttgaccgga	ccaaatcacc	tgggggttgg	acagagaaca	gaagctgtgc	agtaactgca	1380
atgttcccc	caggacggca	gaaactaaaa	agccccaaaga	ggacttcttt	ctcttctgct	1440
gcagatgagt	ggcatcgata	ctaaaagctg	ccccaaacag	gcaaatttta	cctttgtttc	1500
taaagcacca	tcatgtagga	gagccatctg	agggctgggc	aacctcgag	gactcacttt	1560
tgggtggcct	aggctacctt	ggtgttctct	cacataatgt	tcaaggggac	atagtgcaca	1620
agttgaaaag	gctctgttgc	ttttccgtga	gtgcagcagg	gattaaagca	gtctctgctc	1680
cctgtggggc	ctcttgtggg	atatgtctct	cacctaccc	ccacaactca	ggggctcaag	1740
gaccaggcct	cgtgcc					1756

<210> 35

<211> 1545

<212> DNA

<213> Homo sapiens

<400> 35

ggcagagct	cgtgccgctc	gtgccgagag	aagcgaggtt	ctcgttctga	gggacaggct	60
tgagatcggc	tgaagagagc	gggcccaggc	tctgtgagga	ggcaaggag	gtgagaacct	120
tgctctcaga	gggtgactca	agtcaacaca	gggaacctct	cttttctaca	gacacagtgg	180
gtcgcaggat	ctgacaagag	tccaggttct	caggggacag	ggagagcaag	aggtcaagag	240
ctgtgggaca	ccacagagca	gcactgaagg	agaagacctg	cctgtgggtc	cccctcgccc	300
aagtcctgcc	cacactccca	cctgtctacc	tgatcagagt	catcatgcct	cgagctccaa	360
agcgtcagcg	ctgcattgct	gaagaagatc	ttcaatccca	aagtgcagca	caggccctcg	420
agggtgcaca	ggctccccct	gctgtggagg	aggatgcttc	atcatccact	tccaccagct	480
cctcttttcc	atcctctttt	ccctcctcct	cctcttctct	ctcctcctcc	tgctatcctc	540
taataccaag	caccccagag	gagggtttct	ctgatgatga	gacaccaaat	cctccccaga	600
gtgctcagat	agcctgctcc	tccccctcgg	tcgttgcttc	ccttccatta	gatcaatctg	660
atgagggctc	cagcagccaa	aaggaggaga	gtccaagcac	cctacaggtc	ctgccagaca	720

gtgagtccttt	acccagaagt	gagatagatg	aaaagggtgac	tgattttggtg	cagttttctgc	780
tcttcaagta	tcaaatgaag	gagccgatca	caaaggcaga	aatactggag	agtgtcataa	840
aaaattatga	agaccacttc	cctttgttgt	ttagtgaagc	ctccgagtgc	atgctgctgg	900
tctttggcat	tgatgtaaag	gaagtggatc	ccactggcca	ctcctttgtc	cttgtcacct	960
ccctgggcct	cacctatgat	gggatgctga	gtgatgtcca	gagcatgccc	aagaactggca	1020
ttctcatact	tatcctaagc	ataatcttca	tagagggcta	ctgcaccctc	gaggaggtca	1080
tctgggaagc	actgaatatg	atggggctgt	atgatgggat	ggagcacctc	atztatgggg	1140
agcccaggaa	gctgctcacc	caagattggg	tgacaggaaa	ctacctggag	taccggcagg	1200
tgccctggcag	tgatcctgca	cggatgagt	ttctgtgggg	tccaagggtc	catgctgaaa	1260
ttaggaagat	gagtctcctg	aaatttttgg	ccaaggtaaa	tgggagtgat	ccaagatcct	1320
tcccactgtg	gtatgaggag	gctttgaaag	atgaggaaga	gagagcccag	gacagaattg	1380
ccaccacaga	tgatactact	gcatggcca	gtgcaagttc	tagcgctaca	ggtagcttct	1440
cctaccctga	ataaagtaag	acagattctt	cactgtgttt	taaaaggcaa	gtcaaatacc	1500
acatgatttt	actcatatgt	ggaatctaaa	aaaaaaaaaa	aaaaa		1545

<210> 36

<211> 1177

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (24)

<223> n equals a,t,g, or c

<400> 36

cctggtcata	tactcttggc	atancttttt	ttcctttggc	tttgcattggc	tttttcttca	60
gggtactgtct	cgggtatcatt	ctgctaataca	ttgttacaga	atgggtgactt	catttgtgct	120
aacagtacaa	cagcagattt	gggtcagggt	taatctaaag	gttaactttt	ttttctgggtg	180
cttttttggg	ttgatgactg	tctcactttg	actataccca	tgttttgcat	gcaatgactc	240
atgcatgggt	ttcttaacta	gctaataatta	acaatttatt	ccatataaaa	atgggaatttt	300
gcaacatcct	ttaataaggt	gaggggaagca	tgaacctcag	acttctggga	ctattacata	360
gtaagcacat	gaagtagttt	gataataaat	agcagttcta	gtacttcaca	tttcaccctg	420
gtgtgcaatg	cctttttctg	gggggtgggg	gggtgaggga	aacctggtag	tgaatgtgta	480
gttgggggaat	aaagaaaagc	actaaatcct	gccctttttg	tgtgggtttcc	ttttgatata	540
actaggttat	tcataatgta	tacctagaaa	agtgaatttg	aaaataccaa	aagatgtatc	600
atttttattt	gaatccatca	tgcatgtgtac	atttcagata	atttccttca	gtctccagat	660
aggagtgtat	ccaaacatct	aattttatgt	gcactgtgta	tcttatatga	atgttttatt	720
ttatatacca	catgcaaaaa	tgtccatatg	cactatttaa	atgtttttaa	taatataattc	780
cttctttata	atgctaaatc	tatatgagta	ccatattttt	ataagtcagt	gggtctgactg	840
gtttcatttt	agaattaaca	gctgcttcaa	tatgttattc	aatgttaatg	tttggctgtg	900
agtagaatat	gtaaaagtgg	catggcagca	cttatgctct	gtgacagtat	tgtgtgtcat	960
agttgagcag	tagctggtag	aattaggcag	ttgggtgatag	ttttactttg	gtacaaataa	1020
aaactgtata	tctatatata	aataatatat	agatatatat	gtccaccagt	ataatggcat	1080
tgctgtgtct	ggcacttcat	tgtacagact	tttataataa	aagaacttga	aagttctaaa	1140
aaaaaaaaaa	aaaaaaaaaa	aaaaaaaggg	gggggggg			1177

<210> 37

<211> 1173

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (137)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1166)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1168)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1171)

<223> n equals a,t,g, or c

<400> 37

cctgtcctgc	tgtctctgct	gctgcttctg	ggctcctgctg	tccccagga	gaaccaagat	60
ggctcgttact	ctctgacct	tatctacact	gggctgtcca	agcatgttga	agacgtcccc	120
gcgttttcagg	cccttgntca	ctcaatgacc	tccagttctt	tagatacaac	agtaaagaca	180
ggaagtctca	gccccatgga	ctctggagac	aggtggaagg	aatggaggat	tggaagcagg	240
acagccaact	tcagaaggcc	agggaggaca	tctttatgga	gacctgaaa	gacatygtgg	300
agtattacaa	cgacagtaac	gggtctcag	tattgcagg	aaggtttgg	tgtgagatcg	360
agaataacag	aagcagcgga	cattctggaa	atattactat	gatggaaagg	actacattga	420
attcaacaaa	gaaatcccag	cctgggtccc	cttcgaccca	gcagcccaga	taaccaagca	480
gaagtgggag	gcagaaccag	tctacgtgca	gcttgccaa	gcttacctgg	aggaggagt	540
ccctgcgact	ctgcggaaat	acctgaaata	cagcaaaaa	atcctggacc	ggcaagatcc	600
tccctctgtg	gtggtcacca	gccaccaggc	cccaggagaa	aagaagaaac	tgaagtgcct	660
ggcctacgac	ttctacccag	ggaaaattga	tgtgcactgg	actcgggccg	gcgaggtgca	720
ggagcctgag	ttacggggag	atgttcttca	caatggaaat	ggcacttacc	agtcctgggt	780
gggtggtggca	gtgccccgc	aggacacagc	cccctactcc	tgccacgtgc	agcacagcag	840
cctggcccg	cccctcgtgg	tgcctggga	ggccagctag	gaagcaagg	ttggaggcaa	900
tgtgggatct	cagaccag	agctgccctt	cctgcctgat	gtgggagctg	aaccacagaa	960
atcacagtca	atggatccac	aaggcctgag	gagcagtgtg	gggggacaga	caggagggtg	1020
atttgagac	cgaagactgg	gatgcctgtc	ttgagtagac	ttggacccaa	aaaatcatct	1080
caccttgagc	ccacccccac	cccattgtct	aatctgtaga	agctaataaa	taatcatccc	1140
tccttgcccta	gcaaaaaaaa	aaaaangngg	ngg			1173

<210> 38

<211> 1927

<212> DNA

<213> Homo sapiens

<400> 38

ccacgcgtcc	gggcccggact	gaggtcttta	cagtggtccc	tgctggccct	tggtgaccgg	60
tcgcctcagt	tccgaccgg	accgtagcg	tgctgcgctg	acgtggctcc	cggagtagg	120
gctggcgtag	ggccgccatg	ttgcagcagg	atagtaatga	tgacactgaa	gatgtttcac	180
tgtttgatgc	ggaagaggag	acgactaata	gaccaagaaa	agccaaaatc	agacatccag	240
tagcatcggt	tttccactta	ttctttcgag	tcagtgaat	catcgcttat	cttctctgtg	300
agttgtcag	cagcagcttt	attacctgta	tggtgacaat	tatcttgttg	ttgtcgtgtg	360
acttttgggc	agtgaagaat	gtcacaggta	gactaatggt	tggcctacgt	tggtggaatc	420
acattgatga	agatggaaag	agccattggg	tgtttgaatc	tagaaaggag	tcctctcaag	480
agaataaaac	tgtgtcagag	gctgaatcaa	gaatcttttg	gttgggactt	attgcctgtc	540
cagtactgtg	ggtgatattt	gcttttagtg	cactcttctc	cttcagagta	aagtggttgg	600
cgggtggtat	catgggtgtg	gtgctacaag	gtgccaacct	gtatggtaca	tcagggtgaa	660
ggtgcgcagc	agaaagcatt	taaccagcat	ggctacttca	tattttggaa	agcagttttt	720
aagacaaaac	actggagatg	atcagacttc	ctgaatagag	aaagcttatg	tgctttgtta	780
cattggggaa	caactgaaga	gattcttgac	tcaacctttt	agagcttagt	ccatgttgca	840
acgaggagt	ttggctttgt	ttttccactt	aaaaacttta	tttataaaaa	ggaaaagtag	900
ttttcatatt	aagtttttat	ttcctttcca	gcagttgggg	ctagaaagta	tgtgttgcca	960
ctagaaacat	tgtcaagatt	tgttctgtgg	tgtaggtatg	cacattccat	aggtatgcac	1020
acggccatgt	aatatcagta	tatcccaagt	taatgaaagt	gttcattttac	ataggtaatg	1080
gagacctttg	cattttgatc	catagaacat	aggaggatgt	tcttagtctg	tctcaaagct	1140
ctatatgttt	acatatattt	tctgtagatt	gttttcagga	gaaagttttg	cttctatggg	1200
aagagtgagc	actttggcct	atgtataagt	tagaaataat	tgttagtttt	taatatgcac	1260
ttcgtgggga	aattttcttag	acgtatgcaa	gcaagtga	acaattagg	ccagtgggat	1320
taactacttt	ataaaatttt	atttttgttt	gtaagaagtc	atctacttaa	ggcccagtta	1380
atataagtgg	aatcatcata	gtttaaggaa	taccagaga	ttgctgctgt	tctattttatt	1440
ttacagaaag	grtagctaga	ttgaaagctc	ttcagtgagc	cttgagctaa	tagatctttt	1500
accactaaaa	gagcattatt	ctcatgtcat	aatgagaata	ataattttaca	tacttggcat	1560
aataaatgcc	taaaagacat	tttattttct	gaatctat	ttttcttgct	ataatgggga	1620
tattgtaaat	tatgcatttg	tattaatgg	atttcttaaa	gcaatctatg	taactgtaaa	1680
ttaaaccaat	ctacaaacta	ttgtaggcat	ctgtaaatc	tgtgttaggt	attataaaat	1740

ttgttgaagt	cttaatcagc	agattatggt	gtgaatatat	ttgtacattg	ttaaaatagt	1800
ttttaagatt	atttgtttta	ttgaataagt	gtcttattgg	agtgatagct	ttgaagggtc	1860
aaaactttat	atttgtataa	aattctacta	tttacaaggc	aaaaaaaaaa	aaaaaaaaaa	1920
aaaaaaa						1927

<210> 39
 <211> 532
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (467)
 <223> n equals a,t,g, or c

<220>
 <221> SITE
 <222> (513)
 <223> n equals a,t,g, or c

<400> 39						
ggcagagccc	ccacctgcca	gagctgatcc	tccctaggcc	ctgcctaacc	ttgagttggc	60
ccccaatccc	tctggctgca	gaagtccctt	tacccccaat	gagaggaggg	gcaggaccag	120
atctttttgag	agctgagggg	tgagggcatt	gagccaacac	acagatttgt	cgccctctgtc	180
cccgaagaca	cctgcaccct	ccatgcgggc	caagatgggg	aatggaactg	aggaagatta	240
taactttgtc	ttcaagggtg	ctatcgtggt	gcagtggggg	ccctcctggt	gtttgacctg	300
accaagcacc	agacctatgc	tgtggtggag	cgatggctga	aggagctcta	tgaccatgyt	360
gaagccacga	tcgtcgtcat	gctcgtgggt	aacaaaatga	cctyagccag	gccccggaag	420
tgcccatgag	gaggcccgaa	tttcgttgaa	aacaatggat	gttttcntga	gactcagcct	480
ggatttacca	tggtgagtag	ctttgagatt	tcngaagaaa	ttttgcgagt	tt	532

<210> 40
 <211> 1129
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (807)
 <223> n equals a,t,g, or c

<400> 40						
cagctcgwcc	tctgcttcc	tacagcacc	ccacctgcca	gagctgatcc	tccctaggcc	60
ctgcctaacc	ttgagttggc	ccccaatccc	tctggctgca	gaagtccctt	tacccccaat	120
gagaggaggg	gcaggaccag	atctttttgag	agctgagggg	tgagggcatt	gagccaacac	180
acagatttgt	cgccctctgtc	cccgaagaca	cctgcaccct	ccatgcggas	caagatgggg	240
aatggaactg	aggaagatta	taactttgtc	ttcaagggtg	tgctgatcgg	cgaatcaggt	300
gtggggaaga	ccaatctact	ctcccgatcc	acgcgcaatg	agttcagcca	cgacagccgc	360
accaccatcg	gggttgagtt	ctccaccgcg	actgtgatgt	tgggcaccgc	tgctgtcaag	420
gctcagatct	gggacacagc	tggcctggag	cggtaccgag	ccatcacctc	ggcgtactat	480
cgtgggtgcg	tgggggccct	cctgggtgtt	gacctaacca	agcaccagac	ctatgctgtg	540
gtggagcgat	ggctgaagga	gctctatgac	catgctgaag	ccacgatcgt	cgctcatgctc	600
gtgggtaaca	aaagtgaact	cagccaggcc	cggaagtgc	ccactgagga	ggccccgaatg	660
ttcgctgaaa	acaatggact	gctcttcctg	gagacctcag	ccctggactc	taccaatggt	720
gagctagcct	ttgagactgt	cctgaaagaa	atcttttgca	aggtgtccaa	gcagagacag	780
aacagcatcc	ggaccaatgc	catcacntct	ggcagtcccc	aggctggaca	ggagcctggc	840
cctggggaga	agagggcctg	ttgcatcagc	ctctgacctt	ggccagcacc	acctgcccc	900
actggctttt	tgggtcccc	tgccccact	tcagccccag	gacctttcct	tgcccttttg	960
ttccagatat	cagactgttc	cctgttcaca	gcacctcag	ggtcttaagg	tcttcatgcc	1020
ctatcacaaa	tacctctttt	atctgtccac	ccctcacaga	ctaggaccct	caaataaagc	1080
tgtttttatat	caaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa		1129

<210> 41

<211> 553
 <212> DNA
 <213> Homo sapiens

<400> 41
 tctcaccctt cggagacgct cgcccgcacg catagtactt gccgcccagc cagcggcgcg 60
 cgccagccac catgctaggt aacaagcgac tggggctgtc cggactgacc tcgccctgtc 120
 cctgctcgtg tgccctgggtg cgctggccga ggcgtacccc tccaagccgg acaaccggg 180
 cgaggacgca ccagcggagg acatggccag atactactca gcgctgcgac actacatcaa 240
 cctcatcacc aggcagagat atggaaaacg atctagccca gagacactga tttcagacct 300
 cttgatgaga gaaagcacag aaaatgttcc cagaactcgg cttgaagacc ctgcaatgtg 360
 gtgatgggaa atgagacttg ctctctggcc ttttcttatt ttcagcccat atttcatcgt 420
 gtaaaacgag aatccaccca tctaccaat gcatgcagcc actgtgctga attctgcaat 480
 gttttccttt gtcattcattg tatatatgtg tgtttaata aagtatcatg cattcaaaaa 540
 aaaaaaaaaa aaa 553

<210> 42
 <211> 599
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (583)
 <223> n equals a,t,g, or c

<220>
 <221> SITE
 <222> (584)
 <223> n equals a,t,g, or c

<220>
 <221> SITE
 <222> (599)
 <223> n equals a,t,g, or c

<400> 42
 aattcggcac gagtctcacc cctcggagac gctcggcccga cagcatagta cttgccgccc 60
 agccacgccc gcgcgccacc accatgctag gtaacaagcg actggggctg tccggactga 120
 cctcgcctt gtccctgtc gtgtgcctgg gtgcgctggc cgaggcgtag cctccragc 180
 cggacaacc gggcgaggac gcaccagsgg agggacatgg ccagatacta ctcrgcgctg 240
 cgacactaca tcaacctcat caccaggcag agatatggaa aacgatcyag cccagagaca 300
 ctgatttcag acctcttgat gagagaaagc acagaaaatg ttcccagaac tcggcttgaa 360
 gacctgcaa tgtggtgat ggaaatgaga cttgctctct ggccttttcc tattttcagc 420
 ccatatttca tcgtgtaaaa cgagaatcca cccatcctac caatgcatgc agccactgtg 480
 ctgaattctg caatgttttc ctttgtcatc attgtatata tgtgtgttta aataaagtat 540
 catgcattca aaaaaaaaaa aaaaawaaaa aaaaaaaaaa acnngggggg gggccccgn 599

<210> 43
 <211> 1077
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (523)
 <223> n equals a,t,g, or c

<400> 43
 acccagcggt ccgagagtcc accttgcgac cgtatccgct agcgcggcct gggatgcgct 60
 tgggctccct gttcgttccc acatgcaggg cagcacaagg agaatgggcg tcatgactga 120
 tgtccaccgg cgcttctctc agttgctgat gacctatggc gtgctagagg aatgggacgt 180
 gaagcgcttg cagacgcact gctacaaggt ccatgaccgc aatgccaccg tagataagtt 240
 ggaggacttc atcaacaaca ttaacagtgt cttggagtcc ttgtatattg agataaagag 300

aggagtcacg	gaagatgatg	ggagacccat	ttatgcgttg	gtgaatcttg	ctacaacttc	360
aattttccaaa	atgggtacgg	attttgcaga	gaatgaactg	gatttgttta	gaaaggctct	420
ggaactgatt	attgactcag	aaaccggctt	tgcgtcttcc	acaaacatat	tgaacctggg	480
tgatcaactt	aaaggcaaga	agatgaggaa	gaagggaagc	gancagggtg	tgcagaagtt	540
tgttcaaaac	aagtggctga	ttgagaagga	aggggagttc	accctgcacg	gccggggccat	600
cctggagatg	garcaatata	tccgggagac	gtaccccgac	gcgggtgaaga	tctgcaatat	660
ctgtcacagc	ctcctcatcc	aggggtcaaa	ctgcgaaacc	tgtgggatca	ggatgcactt	720
accctgcgtg	gccaagtact	tccagtcgaa	tgctgaaccg	cgctgcccc	actgcaacga	780
ctactggccc	cacgagatcc	caaaagtctt	cgaccctgag	aaggagaggg	agtctgggtg	840
cttgaaatcg	aacaaaaagt	cctgcgggtc	aggcagcatt	agccatcgtg	ccctgctgag	900
gggctggctg	ccttgagtg	cctgatcgcc	acagcccttc	ttggaagaaa	ggcgctcygtg	960
tttcagggtt	cacgcgagtc	acctctttcg	tcttaatgtt	caccgtccac	agctttggaa	1020
taaaccatcc	tgggaagtr	aaaaaaaaa	aaaaaaaaa	tttggggggg	ggggccc	1077

<210> 44

<211> 1904

<212> DNA

<213> Homo sapiens

<400> 44

gctaagctgc	agtgatgttg	cctatatatta	aattttctca	aatggccaag	ctctgatggg	60
ctactttatt	tgagcaatag	ttgagactta	attgcctata	aataaataaa	caaataamct	120
atgtggtttt	ttttctcaca	acatctggcc	tatatgtctt	gtcaggargc	catggctcca	180
atgtaaagta	catagttctt	acatactttc	aactgcagct	ggccccgtgac	ctcaccaggt	240
wtcagagatg	ttctwaaagg	aagccagctg	tggcagggtca	cagattcatg	ggaaatggaa	300
agaaccaagg	aatatagctc	ttgcctcacc	ttctacccca	ctgcagatat	agttcaagcc	360
agagtaatgg	aagaacttaa	cttactagcc	tctcagggtg	ctcctatccc	tacctcccag	420
tgtacagccc	ctccccatct	ctttagtccc	ctttccctca	cttccccctt	tataatgtca	480
cacaaatcag	ggacagtagg	atcacattat	aacctacttt	gtcatagggg	ttcgattttt	540
cttatatcaa	atcatgtttc	ctgaaaccca	gctggggcat	atgcactcaa	tgtctaatac	600
atacttatta	atgtaccgga	tattggcctt	gccccctggat	atcagcaata	tattataaaa	660
ggttccagta	gatgagacga	ttgagtctga	atacaattgc	agtaaattgt	gccaataaag	720
atattgtact	gttacgggtc	tagagttaaa	gccgcttgaa	tgcagcatgc	acattcatgt	780
aaacagacaa	tcagggttagg	cctagaataa	ccacaaaaat	tctattggcc	ttactgcagc	840
cacctatatg	tagaacaatg	gaggagatag	tttgtgggtc	attattgtac	cctgtttcat	900
ccattagcat	cagaatctct	ctttcagggtc	atltattaaa	tatgattgaa	atgtttaaaa	960
gttccctgaac	atgattcatg	atgattaaaa	tatcatacaa	ctgataaaa	actttaagaa	1020
ctttatatat	ttcctgttgc	ctcaaaatgt	aacagaaaat	attcttagag	ctttgatttt	1080
agctatccta	attactgcaa	ataaatatatt	gttctttatag	ttttaaatca	aaaagaaaag	1140
tcttggtata	aaaccttaag	cttgaaatca	tattataaaa	atrtattgta	catagtggaa	1200
aattttcagt	agctaattta	aaatttcaga	aatgtctatt	aaagaatttt	gattcaagta	1260
tttaaactgt	ttagttatgc	atgcttctta	ttaacggaaa	atgataatac	catttagttt	1320
agtgtacagt	atgagaagca	atacctaatc	ctatgttgct	attgtatttt	ttcctagttg	1380
gtgtgcctgc	tcagaaaaac	atatactgta	tgtgtataca	tacctgtgta	tatataaaag	1440
gtcaatttat	atatttttct	ataggaaaa	ggagtaacaa	gttccctatc	tcccatattt	1500
atltgtccat	agtaaaatgg	ccacattgat	gataatttct	agaactagtt	tctgagattg	1560
tcagcccttt	gtctaaaata	atggcagtat	taatgattga	cttctgtcac	tgccatagtt	1620
acctggattg	tcagccttgg	tagcctttgt	ctaaagtcc	aaagagttcc	aaaaaaaatg	1680
tgttgaaaatt	taattgctaa	atagtgggtg	gtgattcttt	acagtaggaa	ttgtaataat	1740
tttcttgcaa	ataagttatt	tactgctatt	gatattgaat	aatttgtctt	ttattcagat	1800
atattttcaa	aagcatgaat	atatgattat	tcataaattg	tatactttac	cagtaagttt	1860
tcagaggaaa	taaagacttt	taaatccttt	tcaaaaaaaa	aaaa		1904

<210> 45

<211> 1625

<212> DNA

<213> Homo sapiens

<400> 45

caagaacaaa	tctgaaggag	gcctctgaca	tcaagcttga	accaaatacg	ttgaatggct	60
ataaaagcag	tgtgacggaa	ccttgccccg	acagtgggtg	acagctgcag	ccagctcctg	120
tgtctgcagga	ggaagaactg	gtcatgaga	ctgcacaaaa	aggggaggca	aagtgtcata	180
agagtgcac	aggcatgtcc	aaaaagaagt	cacgacaagg	aaaacttgtg	aaacagtttg	240
caaaaataga	ggaatctact	ccagtgcacg	attctcctgg	aaaagacgac	gcggtaccag	300

atttgatggg	tccccattct	gaccaggggtg	agcacagtgg	cactgtgggc	gtgcctgtga	360
gctacacaga	ctgtgctcct	tcacccgctg	gttggttcagt	tgtgacatca	gatagcttca	420
gaacaaaaga	cagcttttaga	actgcaaaaa	gtaaaaagaa	gaggcgaatc	acaaggatatg	480
atgcacagtt	aatcctagaa	aataactctg	ggattcccaa	attgactctt	cgtaggcgctc	540
atgatagcag	cagcaaaaaca	aatgaccaag	agaatgatgg	aatgaactct	tccaaaataa	600
gcatcaagtt	aagcaaagac	catgacaacg	ataacaatct	ctatgtagca	aagcttaata	660
atggatttaa	ctcaggatca	ggcagtagtt	ctacaaaatt	aaaaatccag	ctaaaacgag	720
atgaggaaaa	taggggggtct	tatacagagg	ggcttcatga	aaatgggggtg	tgctgcagtg	780
atcctctttc	tctcttggag	tctcgaatgg	agggtgatga	ctatagtcag	tatgaggaag	840
aaagtacaga	tgattcctcc	tcttctgagg	gcgatgaaga	ggaggatgac	tatgatgatg	900
actttgaaga	cgattttatt	cctcttcctc	cagctaagcg	cttgaggtta	atagttggaa	960
aagactctat	agatattgac	atctcttcaa	ggagaagaga	agatcagtct	ttaaggctta	1020
atgcctaagc	tcttgggtctt	aacttgacct	gggataacta	ctttaaagaa	ataaaaaatt	1080
ccagtcaatt	attcctcaac	tgaagtttta	gtggcagcac	ttctattgtc	ccttcactta	1140
tcagcatact	attgtagaaa	gtgtacagca	tactgactca	attcttaagt	ctgatttgtg	1200
caaatttttta	tcgtactttt	taaatagcct	tcttacgtgc	aattctgagt	tagaggtaaa	1260
gccctgttgt	aaaataaagg	ctcaagcaaa	attgtacagt	gatagcaact	ttccacacag	1320
gacgttgaaa	acagtaatgt	ggctacacag	tttttttaac	tgtaagagca	tcagctggct	1380
ctttaatata	tgactaaaca	ataatttaaa	acaaatcata	gtagcagcat	attaagggtt	1440
tctagtatgc	taatatcacc	agcaatgatc	tttggttttt	tgatttattt	gctagatgtt	1500
tcccccttgg	agttttgtca	gtttcacact	gtttgctggc	ccaggtgtac	tgtttgtggc	1560
ctttgttaat	atcgcaaacc	attggttggg	agtcagattg	gtttcttaaa	aaaaaaaaaa	1620
aaaaa						1625

<210> 46

<211> 593

<212> DNA

<213> Homo sapiens

<400> 46

gatgcagttt	gcttggcaga	gctataagcg	ttatgcaatg	gggaaaaacg	aactccgtcc	60
actaacaana	gatggctacg	agggtaacat	gttcggaggc	ctcagcgggg	caacagtcac	120
tgactccctc	gataccctct	acctcatgga	gctgaaggag	gagttccagg	aggccaaggc	180
ctgggtggga	gagagcttcc	acctgaacgt	gagcggagaa	gcatecttgt	ttgagggtgaa	240
catccgctac	atcggggggac	tcctctcagc	cttctacctg	acaggagaag	aggtgttccg	300
aataaaggcc	atcaggctgg	gagagaagct	cctgccggcg	ttcaacaccc	ccacgggaat	360
cccaaagggc	gtggtgagct	tcaaaagtgg	gaactggggc	tgggccacag	ccggcagcag	420
cagcatcttg	gcggagtttg	gatccctgca	cttggaattc	ttacacctca	ctgaactctc	480
tggcaaccag	gtcttcgctg	aaaagggtcag	gaacatccgc	aaggctctca	ggaagwtcga	540
aaagcccttt	ggcctytact	ccaactkagm	catggtgttg	caaacagatc	ccc	593

<210> 47

<211> 1792

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (487)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1306)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (1686)

<223> n equals a,t,g, or c

<400> 47

tttttttttt	tttttttttc	ttattcaata	aactctttta	ttctagtcca	cagatttttc	60
cattacaata	ataatgaaaa	tagcactagt	aatttgtaac	actgaggccc	aaagggaaac	120

ccctcctcaa	attataaggt	aaatgacaca	aagttgaaca	taggggtcagt	gttggggcaaa	180
aagcattttaa	aatatagata	acgggggtcaa	gattttgtgt	gtgtgcaaac	actgggtttt	240
gtttttcagg	atgacaccat	tttagaaagt	gcatgatttt	gaaaactata	tgtgtaattg	300
tgacaaaact	aaactgtaga	gaaaagacaa	aatcaagcaa	aaacaaaaac	caagaaacca	360
aaaggaagca	aatcaaatac	aagggggcgga	tatgcaaacc	tccggtcctt	tggccctgga	420
atggcaagtg	agtgggcttc	atagaaattc	tctgtagaaa	ggaatgtgtg	ggccagtgtg	480
gccatgnccc	tggcctgaca	tcgctgagtc	tccaagtctt	gtctcctcac	agaggaaaag	540
aagttgcttg	tccgacacat	ctgctggggg	tctgcccag	atggggcctt	cgttcccaat	600
ccctttgaga	acagtctcaa	atcctgaaaa	ggcaaggcat	cctgcggcgg	ccccaggggc	660
gcggcaggag	atggggtcag	tgtctgcccc	agctctgccg	gagctgtctg	agtggttcac	720
cgggagtggg	tgggcctcgg	tgttgaacac	ccagtcttcc	agggagagca	agtcactctc	780
agagaacaga	agatagagat	actttagtgt	ctccgctaga	aagaagctct	gctgcttggt	840
gtcgtgggtg	gggggtgctac	tgtacacgtc	ttggatccca	gagaaaccgg	cttctgtccg	900
acagtatttc	tccaaggcca	gcaccacctc	ccagccccac	tccctgtaga	tgggggtgtg	960
ggtctgtcgc	cacaggtaca	tgtagctctc	caccacctct	ggccggagga	tgtagtaret	1020
ctcgtcagc	tgggtggcca	cggcctctct	gccggagtta	aaccagaagc	ctcaggccca	1080
agtttggtgt	ctgagcgggc	gtatgactcg	tgacacgtct	tggtagctcg	ggctgcgagc	1140
tctcggtagt	gggcccctct	ttcttctctg	gcatectcgg	gccaagggcg	atcatgcccc	1200
cggagaaaca	ggccagggtg	cccatcttgt	ggtccagaat	ccccctcgc	cactcggcaa	1260
tgtaggtcag	ccccccggga	garacattca	gcaagtaggt	ctctantcgc	ctccaaggct	1320
tcgtagtaca	tatttttagc	ctccatatct	gtcttgcccc	acatcaacca	ggatttgatc	1380
aaatattcat	aaaaactgtc	cccgaagtct	ccaactgaga	catgggtgtg	caccagttc	1440
ccactcactg	ggctgaggaa	gttggggtag	aggccaaagg	gcttttcgat	cttctctagg	1500
accttgcgga	tgttctgac	cttttcagcg	aagacctggg	tgccagagag	ttcagtggag	1560
tgtaagaatt	ccaagtgcag	ggatccaaac	tccgccaaag	tgctgctgct	gccggctgtg	1620
gcccagcccc	agttcccact	tttgaagctc	accacgccct	ttgggattcc	cgtgggggtg	1680
ttgaangccg	gcaggagctt	ctctcccagc	ctgatggcct	ttattcgga	cacctcttct	1740
cctgtcaggt	agaaggctga	gaggagtccc	ccgatgtagc	tcgtgccgaa	tt	1792

<210> 48

<211> 785

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (704)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (746)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (753)

<223> n equals a,t,g, or c

<400> 48

gcggctgggt	tcttggtgag	cccgggtccc	tcaaggccgg	aaagaaagtc	gggcttctct	60
agccccctgga	ggactcgact	cactgggtgcg	cgatttaggt	ccggagaggc	gttgtgaggt	120
gagctttttc	agaagcgcg	tcccaggaca	cgctcggaag	caagcatccc	cagagctgct	180
tggaaagagg	accaaagacg	tctaaaaagt	catttggaag	tatctctaaa	tatttggtac	240
catgtataag	ctgctaaaga	gaaattgggc	ccaacaaaac	taattgaata	attgaggcag	300
atttgtgtgt	atcatcaaat	tctatccaga	agttgaagaa	tctgaattta	aagattgtgt	360
gcatttaata	agaggatgac	ctttcagttt	aatttcacta	tagaagacca	tctggaaaaat	420
gaattaacac	ccattagaga	tggagctttg	accctggatt	cctcaaaaga	gctgtcagtc	480
tcagaaagtc	aaaaaggaga	agagagggac	agaaaatgtt	ctgcagaaca	atttgacttg	540
cctcaggatc	acttgtggga	acataagtca	atggaaaatg	cagctccctc	tcaagacaca	600
gacagtccac	tcagtgcagc	cagcagttca	aggaacttgg	gagccacatg	ggaaaacagc	660
cctccttgag	agctggccaa	aggrgcmtgc	tatgccttaa	aggntttaaa	gaagrtgttt	720
aggaaaatwa	aagtycttag	gaaacnttta	ccnggggttt	ccmgyctgtt	taagttwttc	780
rgtta						785

<210> 49
 <211> 1433
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (893)
 <223> n equals a,t,g, or c

<220>
 <221> SITE
 <222> (947)
 <223> n equals a,t,g, or c

<400> 49
 gcggaggcct ccgtagtgat ctggccttta ctttctcccc gagtcacggg aagccctcgt 60
 tgacctcaca ggggtggacac ccggaggcga gatcccgttc cgcggagcag agccctttct 120
 catggaacag gacgtgtcgg ggccgctgct ggggaaagca gccgggcccc cagatgctgg 180
 agcgggagca ggccccgggc ccccgagac cctccgcggc accgcccgct cttgtgcctt 240
 tcccggcgtg gtcaccgcc tcaccatctc ggggtgtctt taggagaatc cttcatgcag 300
 ctgcagcagc gtctcttgag agagaaggag gccaagatca ggaaggcctt ggacaggctt 360
 cgcaagaaga ggcacctgct ccgcccggcag cggacgaggg gggagtccccc cgtgatctcc 420
 gtgggtgggg acaccaactg cggaaagacc acgctgatca aggcactgac gggcgatgcc 480
 gccatccagc caccgggacca gctgtttgccc acgctggagc tcacggccca cgccgggacg 540
 ctgccctcac gcatgaccgt cctgtacgtg gacaccatcg gcttctcttc ccagctgccg 600
 caccggcctca tcgagtcctt ctcggccacc ctggaagacg tggcccaactc ggatctcatc 660
 ttgcacgtga gggacgtcag ccaccccagag gcggagctcc agaaatgcag cgttctgtcc 720
 acgctgcgtg gcctgcagct gcccgccccg ctcttgact ccattggtgga ggttcacaac 780
 aaggtggacc tcgtgccccg gtacagcccc acggaaccga acgtcgtgcc cgtgtctgcc 840
 ctgcggggcc acgggctcca ggagctgaaa ctgagctcga tgcggcggtt ttnaaggcga 900
 cggggagaca gatcctcact ctccgtgtga ggctcgcagg ggmgcantca gctggctgta 960
 taaggaggcc acagttcagg aggtggacgt gatccctgag gacggggcgg ccgacgtgag 1020
 ggtcatcatc agcaactcag cctacggcaa attccggaag ctctttccag gatgaacgga 1080
 cgcccacaga ggccctgcgg gtgggggcat cgctgcctgg ggagctgagg cgttaccgct 1140
 gtgttggggg cagcttggtg tcaggtgcag cagggtccctc cttgtctggt tctgcacccg 1200
 tctcgctccc agccatttgc tgggatgacc gtgcaggccc gtgacacggc cgcacctgcc 1260
 ccaaagcggg ccgcccggag gtccactcca agcctgagca tccacacaat tccagtgggc 1320
 cctcggtgcc tgctgtgaac tgctttccct cggaatgttt ccgtaacagg acattaaacc 1380
 tttgwtttta cttccgtgaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa ggg 1433

<210> 50
 <211> 652
 <212> DNA
 <213> Homo sapiens

<400> 50
 ctttctcacc actctcctgc tagccatctc tttggcacta aggccttggg caaattggat 60
 ttctttcatt tttccacact tcaaagaccc atgtttctagg tattctccat agggatagtc 120
 tctttggcat ttatttggtt tttctacgtt ttcagtccca tttactccaa gactcactcc 180
 ctgccaccta gtgcatcaga tacagctact tctggctgac ttttcaaggg ggaccaccct 240
 acctgtcatc tcttcaactgt tcagaaatga ctgtgtcagt ggcacctcaa actcccttgc 300
 tgtccttttc caaggagaca gctaagggtg atggagatgc agaattggacc tcacgttcgc 360
 cctagtcagg actgataccc tttccgtttc agaggattgc caagaaaaaa ctcacagtgtg 420
 aggcagggtg ctctgaggtc ggctgcgggtg tgggaggcac gsctgggcmr gctctctggg 480
 ctggagcagg tggattcgaa ggcctgtcta gcacgagggc ccaaaggctt tgtcagtggc 540
 cagtagctct gccgcctttc ccagagaggg ggtccagggg acatcctgga aggctggggc 600
 ctggggccacc ttctgtcttt gcaagctaga gccagcccaa tagggggcgg at 652

<210> 51
 <211> 2541
 <212> DNA
 <213> Homo sapiens

<400> 51

ccacgcgtcc	ggttttatatt	tttttcacct	tttaaaca	aaattcttta	aaatatattta	60
atgcattctt	ttgaagaggt	agatgttttg	tacattttat	ggctcccaga	gcataatattc	120
agttggtgca	tggtgtggaa	gggggaattg	gaaattaaat	gaaaacctat	gactttgggtc	180
atgtcaatct	gtaagacaca	tcagtaaaag	gggtattatgc	tctgttggtt	ttgtttttttt	240
gttttgcttt	tttttttttt	ttttcttttt	tgtttttttg	tgatgtggct	taaatgcaat	300
agtttctttt	tgggacatat	ttctgccaat	taaagactag	aaggcacaac	ttttttttttt	360
aattaccata	gagaagatac	attaaaaaaa	atcttctgat	gttttgtagc	cataactaaa	420
ttatggtaaa	aatgtgcaact	attgtgaaaa	ggagcaacgt	agttttgggt	tttttggtgt	480
ttgtttgttt	tgctttgttt	tttaagagat	taaaatgttt	ctggataagg	attagcttct	540
cgaagtgtcc	atcattctgt	gtagaagctt	aaatatgtaa	tgtaaccaa	ctccagtatt	600
aaaaatctct	catgttggtt	tctttatata	aagcaagata	acggcatata	acactgccat	660
tacatggcaa	aatgtttgct	accttagttt	aaaaaacaat	ctcaaacaaa	agacttgctt	720
caaggtgttt	ttaaatagca	gtgattcaga	atttttttta	atgaaagtat	aattgcacta	780
accttcttcc	tgctgtctcg	attctgcatt	tggtgtactt	gtgactacgt	tttttcaaat	840
atagatagat	ttaaagtgtt	aatttttttt	tttttagtaa	tcactactat	atcatgtctt	900
ttactctgtt	tataatatca	agtattttct	taaagatata	gatattaaac	cttggtctca	960
tgcaacttag	agtaacatat	acagacaaat	gattgcatga	ggccatgttt	atatgtgtga	1020
ctaataaggc	ttgtcatgat	taacataatc	caggtatgtc	atttctgaag	agaatagtca	1080
tcaaattttt	atctcgaaga	ttttaattaa	ggaattgctt	attgttgagc	ttagcaaat	1140
aataacacta	tttctgtcac	taattatttt	gaggcctttt	agtactaaaa	ttttaacctg	1200
tgttctaagt	agaaactgat	ttaaccacag	taatgcagct	ttgattgatt	tcagcattcg	1260
ttgctttgct	attttttacaa	aacagcattg	attgaagcaa	gtcttggttt	tactaaggta	1320
gggtagcatt	tgctattggt	aaagagaata	aatacactta	atttcacaa	acattgttat	1380
atgtacccca	gttgttggtt	gtggggacta	tgatactgta	ataatatttt	taaaaattta	1440
catcaagaga	ggcagtcatt	cacgatgggt	ttgtgccagc	tctttttagg	gttttggtatc	1500
acattagaga	tatttagaac	atattaccct	gtgacttacg	taggaaacct	aatatgctga	1560
gtatctggca	cttgaattcc	tgctttttat	gctggagggtc	cacatctgtg	gttgacctct	1620
gttattgttt	aaaaaaaaata	ataaaaaatt	aaaaaaaaatc	gtgcaataat	tttaaaatgt	1680
gctcccagga	atagacacaa	atgtttttgag	tatcttttaa	gctgcatttt	ccttttagtga	1740
tgcatattgtc	aatgtgcactg	aatttaaatc	tgaaagtcag	aggtgattat	tgatagtact	1800
tttgtatttt	gatattggaca	gtttattcat	ttgcatacag	ttattgactt	tttcccagct	1860
gattaaaaga	tagtcaagaa	attctgcaat	atagctgcca	aaatagacag	ctacattttt	1920
atgatattgt	catcttttct	gttttttttt	tctttttttt	ctttagctat	tttacttaag	1980
cataatagcc	acaataggac	atataaaaga	ttataaatac	agagctttat	tatcctgacg	2040
tcttggtctt	tttaagtata	tacttttctg	aaaggatatc	attttgtagg	cttggtgtct	2100
tcattgacat	acgattgttt	atttttgctg	ctgttctcaa	catcatcatt	gcctgctgat	2160
gtgccagat	gctgctccaa	tagacagcaa	taagattgtc	tctaatttga	gcagtaacat	2220
gattgcaaga	gaccaagtgt	cacagcttgt	aaagtctgtt	atttggtgatt	cttgcttatt	2280
tttccgcctg	tggttttctg	agaacttatt	cctgatgatc	aattgaaatc	agtagttttt	2340
ctatgctatt	tggtgttgta	taagctactg	taagaaactt	atcataagga	aaaatagaaa	2400
ggaaaacttg	aatcaatact	cattgattaa	aatggaataa	agaaagagca	gctgccactt	2460
ttaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	2520
aaaaaaaaaa	aaaaaaaaaa	a				2541

<210> 52

<211> 701

<212> DNA

<213> Homo sapiens

<400> 52

aacattacaa	gggctttttat	aaaaaacctt	ttgttcatat	ttcttccctt	taaaatatgt	60
aatgtcaaaa	atgactcacc	ttttaaaaa	tatgcatgaa	aacaggtggg	aaacattcag	120
taatacgcta	tttctccaac	atcaagacaa	ctaaaacaaa	tgataaaaa	gtttattttt	180
acactccagc	atatcggttg	agtttttagg	atgtgtatga	atattttaa	cttttaattt	240
cagttttaat	gaaagctgaa	cttaatagg	aaagctagct	cttggttaact	agcaatgatc	300
aggcattggt	tgctctgttc	aggttttctt	atctgtttta	ggtagatttt	ttcagattct	360
gattgtttga	gttaatgggt	gaatttttaa	agtttttagt	tactttaaa	akgattttta	420
atrrcatatt	aatttagaaa	attcctgtgt	ttacttatat	tttaaatgt	gaaattggatc	480
caatcattag	aacagagaga	atagttcttt	gaaactgaaa	tactttagtt	ttactgacct	540
tgtgtaaaga	taatatgaag	aaccagcttc	caaaagaaac	cagcatatgg	cactataaac	600
tatttcattt	gagcaccatt	ctttaccatg	gatataatga	ttatgtatta	tagtggagtg	660
atcatacagk	tcccccaaat	gtgatgggtc	aagggaattt	a		701

<210> 53
 <211> 375
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (153)
 <223> n equals a,t,g, or c

<220>
 <221> SITE
 <222> (313)
 <223> n equals a,t,g, or c

<400> 53
 tgcctctgtg tgtgtgcaag acagagagat aggcattttg tcaagtcage tagttgccta 60
 ggtatctttg tctcacatct ggctgtttcc tcctagagaa ccatccagtt ggctttccag 120
 gtctggaggt gagctaattg atgagtgaat atnagcagtg ggtgttcctc atctctttga 180
 ggattttgct cagagttcac taccaagggg tttctggaac tagggwccat tctttacatc 240
 agttcttgag ggttctttga tatcaggggc aaaatgatcc cttctctttt ctttcttata 300
 tctgtgctt tgnctcctgg gtgattttct tccaagtcag ttgtgggagg tgcctaggaa 360
 caacgctaac acggg 375

<210> 54
 <211> 1146
 <212> DNA
 <213> Homo sapiens

<400> 54
 tcgacccacg cgtccgatag agacgctgaa ccatgaacat tatgagtga aatacgtaat 60
 taatgcgatc cctccgacct tgactgccaa gattcacttc agaccagagc ttccagcaga 120
 gagaaaccag ttaattcagc ggcttccaat gggagctgtc attaagtga tgatgtatta 180
 caaggaggcc ttctggaaga agaaggatta ctgtggctgc atgacattg aagatgaaga 240
 tgcctcaatt tcaataacct tggatgacac caagccagat gggtcactgc ctgccatcat 300
 gggcttcatt cttgcccgga aagctgatcg aattgttaag ctacataagg aaataaggaa 360
 gaagaaaatc tgtgagctct atgccaaagt gctgggatcc caagaagctt tacatccagt 420
 gcattatgaa gagaagaact ggtgtgagga gcagtactct gggggctgct acacggccta 480
 cttccctcct gggatcatga ctcaatatgg aaggggtgatt cgtcaaccgc tgggcaggat 540
 tttctttgcg ggcacagaga ctgccacaaa gtggagcgcc tacatggaag gggcagttga 600
 ggctggagaa cgagcagcta gggaggtctt aaatggcttc ggggaaggtga ccgagaaaga 660
 tatctgggta caagaacctg aatcaaaggga cgttccagcg gttagaatca cccacacctt 720
 ctgggaaagg aacctgccct ctgtttctgg cctgctgaag atcattggat ttccacatc 780
 agtaactgcc ctggggtttg tgctgtacaa atacaagctc ctgccacggt cttgaagttc 840
 tgttcttatg ctctctgctc actggttttc aataccacca agaggaaaat attgacaagt 900
 ttaaaggctg tgtcattggg ccatgtttta gtgtactgga tttactacc tttggcttaa 960
 ttccaatcat tgttaaagta aaaacaattc aaagaatcac ctaattaatt tcagtaagat 1020
 caagctccat cttattttgc agtgtagatc aactcatgtt aattgataga ataaagcctt 1080
 gtgatcactt tctgaaattc acaaagttta acgtgatgtg ctcatcagaa aaaaaaaaaa 1140
 aaaaaa 1146

<210> 55
 <211> 2299
 <212> DNA
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (179)
 <223> n equals a,t,g, or c

<220>
 <221> SITE

<222> (180)

<223> n equals a,t,g, or c

<400> 55

tcagacagtt	tgaatacttg	aatcatgcag	gccaatatta	taatgtgaaa	aggatatctac	60
tctatttaca	ctcccaata	gcgccatata	tgctaaaccg	tagagaatga	gctcgcttgt	120
gtctattcat	catgttttagc	ctttgggattc	tttttttttt	ttccttctat	tctctcccn	180
ccccccccc	cgccctttt	tttytytyt	gcaaaacccat	tttttgggct	gataacgtat	240
gagcttttcc	ctttgcactg	aatgatgttc	tctccgtctc	atcggcagta	tggggggcag	300
ctgtcccagt	gtcaatgttt	actcaagggg	gttcttagga	ggcgtgcgct	ctctactatg	360
ccttgatgtt	gcctacctta	ttgtgggtatc	gtggagttta	aaagatcaag	ttaggatgct	420
gacttaggat	tattaatgaa	agtgttgcac	cagttttttc	atgttgtaaa	actaaagaat	480
ttcgtctctg	agtttgaaaa	actgtggcca	cagctgtgac	ttgcagccca	cctgccaccc	540
aggacggggc	ctgcactttg	aataggcttt	ccattttgtt	ttggagggtc	tcactttgaa	600
ccttcttgtt	tacagatttt	tttgtttgtt	ttttgagaaa	aaaaaatgtt	tactcttcca	660
tcatttaaaa	aaaatgtaaa	agacaaaaaa	aaaatggagg	atgattttaa	agatgctttc	720
tatctctggg	aaaaaggagc	agcattttggc	catgttcttt	tgtttttcta	ttcctgtccc	780
aatcaaaga	gcatggttct	caggaaaacc	agttcccag	tttaaaaaaa	aaaaaaaaaa	840
ttccttgtag	tttcttagag	gaaaaaaaga	aaaaccccaa	cttttagcac	tgatactaca	900
tattgctctg	ttaaagaatt	ttctctgcca	aaaaaaaaga	aaaaacaaaa	aaacgcttaa	960
agctggagtt	tgacattctg	ctttcagatg	ctgtcttttt	attagttagt	gatgatgggt	1020
tgctaataat	caataggtaa	taattttttg	taatcccatc	aagtggctcc	atatgtttct	1080
gctctctcgt	gactgtgtta	atgtttaact	gttgtacctt	aaagccgaaa	tcagtaacta	1140
tgcatactgt	aaccaaggta	ttgggcttac	agagttgttt	gttgataaaa	gaaaatttta	1200
aatgttggtg	caaactaacg	agttacacca	ttttaaactt	tctttcctcc	cccccttttt	1260
tgcccacaaa	tggtattata	atgcttgctt	agtcaaagaa	gagagactaa	acaagggtaa	1320
aaattttaac	agtacagaat	ttgccatcat	atcattgcct	tgattctaac	tgtttgtgtc	1380
ctaagatgca	aaagaagtca	gtggctttta	actgtttaca	aatagaatgt	gattgtaaaa	1440
tgtacagttt	ggttgtgttt	gaattatgaa	atctcttcag	atataataaa	ccatgacttt	1500
ttggctgctc	aacattaatt	gtctcctttt	tgtgaattta	tttgtaggct	cttttttata	1560
atgaaagttt	caaagttgct	atgtatgagg	gttctcatag	agcaaccgat	taaaaatcta	1620
agcaaatatt	tgaacatttt	atctgaactc	atcacaattt	caccctgaaa	taatgtgaga	1680
acaatgggaa	actgtagctt	gtccttccc	accctctctg	agcatctttg	ggatcttgtt	1740
gctcaaaact	cttctgtgac	ttcatcttcc	ccaccatttg	tgcccatctc	aagcctcagc	1800
aagaaacccat	gtggaacatg	aagcttaatg	acttgacagt	gtactagtgt	taaactctca	1860
tacctctgtt	acaaagcgag	aaacgccaca	cccgactgg	ccttttcttc	ccccctcacg	1920
gccctcgctt	ctccctgcag	gagctcgggg	gcgaaacctg	tgtatggatt	tcagtgtatg	1980
acttcagatc	atgtccaac	ttgccagggtg	tgagctaagt	ttgtcggaca	ccttactata	2040
agcaaatgtt	attcagtgcg	ttcaatgtat	attgacttcc	atactgggtt	ttccaaaaac	2100
caaaggtagc	tttgaaaaac	catgtctgga	aatgtttgga	gcgttaagct	gattgacctt	2160
ctgaccttgg	ggctttgagt	agtatataat	tcataactgc	gttaattgta	ttgttaaagt	2220
gtttggggagt	tttttgcgct	tgttatgtgg	aaataaagtg	tttgatttaa	aaaaaaaaaa	2280
aaaaaaaaaa	aaaaaaaaaa					2299

<210> 56

<211> 2259

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (2213)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (2242)

<223> n equals a,t,g, or c

<220>

<221> SITE

<222> (2247)

<223> n equals a,t,g, or c

<220>

<221> SITE
<222> (2250)
<223> n equals a,t,g, or c

<220>
<221> SITE
<222> (2253)
<223> n equals a,t,g, or c

<400> 56
ccgcagccca tctgctggca tcaktacctg gtgttggggac agcaggatag gkttctaaag 60
gtgggttttyt atccaaacga ccaaaaaacc aacagtaaca ccagtgaac cccacactgt 120
cgggcttata aaaatctgtg ccatcatggt gattttatcc aagactgctc cacttacccc 180
agtgcagggg acaagtttct gttgaaactt tagatagcag aattatttgc aattttagtc 240
atagaaaaga tttttaaat tttttacaaa aggtttttta acagattagg gtagggtgatg 300
gtttaaatca attaagtggc attggaacc taggggttcc ttttgattaa gagccttttt 360
tgtttctgct ctttgtcagc tttcagggga gaaggaggcc actggaaaat tatttcctta 420
agtgcaggct gttgactgcg tatgccaaaa agggacagga ggcatgggat agcagggtctg 480
gtgacacagc tagggctctc ctagcagctc ctccctctcc ctcccaaggc cccaggaat 540
cccttctctc catgtcctgg cagcaggacc ccaggctaca tatggaagg agagatgtgg 600
gggtcctgtr tectggagta ttatgtctcc ccacctctg cagttttctc tgaacatgta 660
tggtgcccac ggtgggagcg tggctactgt gcagttgtgc acagatgtct ttcctttacc 720
gttggccttt ctgtctgcct ctcttctctc tctgcagccc aaatggaaaa caattattta 780
ctccattgga gggaaaggaa gagtcttaga attcctaagg gaaccttagc ataaagggtt 840
tggggaagga ggccgtaggc scggaggaa gcaattccac ttggtttgac aacttctgcc 900
actcccatgt cagtgactt gcacttctta aagagattgc tttataacac taagacatcc 960
tttctaaaga ttcaagtgga cttgactaag ctgagggtcc acgaaataga atatgacatg 1020
tgagctgttt ttggaaaacg aagatggaga gagcacttcc ccgtaacgaa agcaaagtgg 1080
taagcacagg gtgagaccct tttacacaga atggtggaga gaaaagagaa tgctgaaaag 1140
tggctcagat gcagagtgtt ctgtggagaa actgcagccc cacttctgtt tccctggagt 1200
ctcccaatgg atcattcagg agtgcctat gtgagaattg agccaaggaa aatactcatg 1260
caaccagcct gagtcgcggt gaggggacga gaggtgtgac acacattggt agttattttg 1320
caccagcagt gcctttctca ctgggggtac ttggaccctc agatcttctt ttctaatagc 1380
catttgccac cccaagtgg atgtcgccca tttctcttta aaacaccttc cctacctttc 1440
ccatgtactc agtttagctc tcaaagaagg ggtgaatcat aaagccagtg aaaatttcac 1500
cctctgaggg agttcccca tctgaagggg aagaggggtga cctcagcggc ttttctccca 1560
aaaatcggct gaagctggt tgtggatcct tgttctctc ctgaccccat ctggctgctg 1620
ccccgtctcc caccctgtc cccggggctc gctggccctg cactccgcct tagtctggg 1680
gccggcgaca cagtgggggc tctcacttg ctgcagtgtc atagcaataa aatgtgattc 1740
ttgggggtccc cccaggagc tgcccattgg tttatttatg aacctggtt tccgggagtc 1800
ggggaggaga tgactttgct tctgtgcaca gccccgtctt ccaggagcca cgactcagaa 1860
gaaaaggggt ctcagacttt tgttatacac atttgctttg tgtaataaaa tgtttacaat 1920
tttatatgaa agatggaata agcgctagag cttccaactg tatatttttt acttttatag 1980
attttaaaac tatgactctt tatatgtgtg ttttggggga gctatgataa gttttatggc 2040
aaacggttgg tattgttaac tttttattgt catcaaaagt tcataaaagt cctattaatc 2100
cccatattct tctactgcc ttaactctgg tatacaccaa aaagaaatct ttactttcct 2160
tgttttatca ttataaaaat aaagtatttt gctagtatgg aaaaaacctt tgnatttgac 2220
gtcacctggg gtctgctggc anaaagnttn ggngaattg 2259

<210> 57
<211> 1325
<212> DNA
<213> Homo sapiens

<220>
<221> SITE
<222> (1302)
<223> n equals a,t,g, or c

<220>
<221> SITE
<222> (1313)
<223> n equals a,t,g, or c

<400> 57

ttaaaaacaag	atacatacat	agtataaacac	acctcacagt	gttaagattt	atattgtgaa	60
atgagacacc	ctaccttcaa	ttgttcatca	gtgggtaaaa	caaattctga	tgtacattca	120
ggacaaatga	ttagccctaa	atgaaactgt	aataatttca	gtggaaactc	aatctgtttt	180
taccttttaa	cagtgaattt	tacatgaatg	aatgggttct	tcactttttt	tttagtatga	240
gaaaattata	cagtgcctaa	ttttcagaga	ttctttccat	atgttactaa	aaaatgtttt	300
gttcagccta	acatactgag	ttttttttaa	ctttctaaat	tattgaattt	ccatcatgca	360
ttcatccaaa	attaaggcag	actgtttgga	ttcttccagt	ggccagatga	gctaaattaa	420
atcacaaaag	cagatgcttt	tgtatgatct	ccaaattgcc	aactttaagg	aaatattctc	480
ttgaaattgt	ctttaaagat	cttttgcagc	tttgcagata	cccagactga	gctggaactg	540
gaatttgtct	tcctattgac	tctacttctt	taaaagcggc	tgcccattac	attcctcagc	600
tgtccttgca	gttaggtgta	catgtgactg	agtgttgggc	agtgagatga	agtctcctca	660
aaggaaggca	gcatgtgtcc	tttttcatcc	cttcactctg	ctgctgggat	tgtggatata	720
acaggagccc	tggcagctgt	ctccagagga	tcaaagccac	acccaaagag	taaggcagat	780
tagagaccag	aaagaccttg	actacttccc	tacttccact	gctttttcct	gcattkaagc	840
cattgtaaat	ctgggtgtgt	tacatgaagt	gaaaattaat	tctttctgcc	cttcagttct	900
ttatcctgat	accatttaac	actgtctgaa	ttaaactgac	tgcaataatt	ctttcttttg	960
aaagctttta	aaggataatg	tgcaattcac	attaaaattg	attttccatt	gtcaattagt	1020
tatactcatt	ttcctgcctt	gatctttcat	tagatatatt	gtatctgctt	ggaatatatt	1080
atcttctttt	taactgtgta	attggtaatt	actaaaactc	tgtaatctcc	aaaatattgc	1140
tatcaaatta	cacaccatgt	tttctatcat	tctcatagat	ctgccttata	aacattttaa	1200
taaaaagtac	tattttaatga	ttaaaaaaa	aaaaaaaaa	aaaaaaaaa	aaaaaaaaa	1260
aaaaaaaaag	gaaaaaaaa	aaaaaaaaa	aaaaaaaaa	angggggggg	ggnccaaaa	1320
aaaaa						1325

<210> 58
 <211> 832
 <212> DNA
 <213> Homo sapiens

<400> 58						
gcgtcgacat	agaattgaag	ttgctcgta	gctgattgaa	gataaggaga	ttggcctgga	60
ttatccaggt	aggctcaatg	taatcaggaa	gggcctttta	agtgagagag	ggasgsagaa	120
gaggaagtca	gagcgatgtg	ctgtgaaatc	tactaccgtt	tgctgggttt	gaaaatggag	180
aaaaagagt	aggaactgag	aaacatggat	ggccttggga	acgtggaaaa	gggtcactga	240
aatgggacga	catgaactca	aggaggctat	ttatgaccat	gtcatttgca	acatgaagaa	300
agcttatctg	gagtgaagt	aaatgagacc	aacagagatr	agagaccg	agaaatcctg	360
gttacactgc	ttgaatcctg	tcagtcctat	actggagtcc	tgtaataaca	aaataatagt	420
aataatccct	ctgtttctta	tgtttatgcc	aacttcaaca	aaaagaaact	tgactaagag	480
acaatataag	aayttaatgt	gtaattaaga	aagaactctc	caccacgggg	aatgtgaaag	540
gtatatgagt	cccttttcac	gatgcgatgt	catgtctttt	aaataagcca	tactttatgt	600
tcaataaaaa	gagaataagc	aggattcgcm	agagaacaca	atcccttttt	aactgctggg	660
aagatacytt	tagtcattaa	tgrctggacg	acaatttggg	rcacmtatat	ggatattggc	720
cggtttgta	tgatgtgatt	gggcctctaa	gtgacaacat	tgttccctgt	atagagtga	780
tggcaagtgc	atttataaaa	ttggccatca	tggctgttaa	atttataaaa	aa	832

<210> 59
 <211> 132
 <212> DNA
 <213> Homo sapiens

<400> 59						
cttgagcccc	tgagttgtgg	gggtagggtg	aagagcatat	cccacaagag	gccccacagg	60
gagcagagac	tgctttaatc	cctgctgaca	tcacggaaaa	gcaacagagc	cttttcaact	120
ttgtcactat	gt					132

<210> 60
 <211> 54
 <212> PRT
 <213> Homo sapiens

<400> 60	
Met Ala Phe Leu Gln Val Leu Ser Arg Tyr His Ser Ala Asn His Cys	
1 5 10 15	

Tyr Arg Met Val Thr Ser Phe Val Leu Thr Val Gln Gln Gln Ile Trp
20 25 30

Val Arg Leu Asn Leu Ser Val Asn Phe Phe Phe Trp Cys Phe Phe Gly
35 40 45

Leu Met Thr Val Ser Leu
50

<210> 61

<211> 117

<212> PRT

<213> Homo sapiens

<400> 61

Met Phe Pro Phe Tyr Ser Cys Trp Arg Thr Gly Leu Leu Leu Leu Leu
1 5 10 15

Leu Ala Val Ala Val Arg Glu Ser Trp Gln Thr Glu Glu Lys Thr Cys
20 25 30

Asp Leu Val Gly Glu Lys Gly Lys Glu Ser Glu Lys Glu Leu Ala Leu
35 40 45

Val Lys Arg Leu Lys Pro Leu Phe Asn Lys Ser Phe Glu Ser Thr Val
50 55 60

Gly Gln Gly Ser Asp Thr Tyr Ile Tyr Ile Phe Arg Val Cys Arg Glu
65 70 75 80

Ala Gly Asn His Thr Ser Gly Ala Gly Leu Val Gln Ile Asn Lys Ser
85 90 95

Asn Gly Lys Glu Thr Val Val Gly Arg Leu Asn Glu Thr His Ile Phe
100 105 110

Asn Gly Arg Gln Phe
115

<210> 62

<211> 183

<212> PRT

<213> Homo sapiens

<400> 62

Met Val Arg Met Val Pro Val Leu Leu Ser Leu Leu Leu Leu Leu Gly
1 5 10 15

Pro Ala Val Pro Gln Glu Asn Gln Asp Gly Arg Tyr Ser Leu Thr Tyr
20 25 30

Ile Tyr Thr Gly Leu Ser Lys His Val Glu Asp Val Pro Ala Phe Gln
35 40 45

Ala Leu Gly Ser Leu Asn Asp Leu Gln Phe Phe Arg Tyr Asn Ser Lys
50 55 60

Asp Arg Lys Ser Gln Pro Met Gly Leu Trp Arg Gln Val Glu Gly Met
65 70 75 80

Glu Asp Trp Lys Gln Asp Ser Gln Leu Gln Lys Ala Arg Glu Asp Ile
85 90 95

Phe Met Glu Thr Leu Lys Asp Ile Val Glu Tyr Tyr Asn Asp Ser Asn
 100 105 110
 Gly Ser His Val Leu Gln Gly Arg Phe Gly Cys Glu Ile Glu Asn Asn
 115 120 125
 Arg Ser Ser Gly Ala Phe Trp Lys Tyr Tyr Tyr Asp Gly Lys Asp Tyr
 130 135 140
 Ile Glu Phe Asn Lys Glu Ile Pro Ala Trp Val Pro Phe Asp Pro Ala
 145 150 155 160
 Ala Pro Tyr Ser Cys His Val Gln His Ser Ser Leu Ala Gln Pro Leu
 165 170 175
 Val Val Pro Trp Glu Ala Ser
 180

<210> 63
 <211> 205
 <212> PRT
 <213> Homo sapiens

<400> 63
 Met Leu Gln Gln Asp Ser Asn Asp Asp Thr Glu Asp Val Ser Leu Phe
 1 5 10 15
 Asp Ala Glu Glu Glu Thr Thr Asn Arg Pro Arg Lys Ala Lys Ile Arg
 20 25 30
 His Pro Val Ala Ser Phe Phe His Leu Phe Phe Arg Val Ser Ala Ile
 35 40 45
 Ile Val Tyr Leu Leu Cys Glu Leu Leu Ser Ser Ser Phe Ile Thr Cys
 50 55 60
 Met Val Thr Ile Ile Leu Leu Leu Ser Cys Asp Phe Trp Ala Val Lys
 65 70 75 80
 Asn Val Thr Gly Arg Leu Met Val Gly Leu Arg Trp Trp Asn His Ile
 85 90 95
 Asp Glu Asp Gly Lys Ser His Trp Val Phe Glu Ser Arg Lys Glu Ser
 100 105 110
 Ser Gln Glu Asn Lys Thr Val Ser Glu Ala Glu Ser Arg Ile Phe Trp
 115 120 125
 Leu Gly Leu Ile Ala Cys Pro Val Leu Trp Val Ile Phe Ala Phe Ser
 130 135 140
 Ala Leu Phe Ser Phe Arg Val Lys Trp Leu Ala Val Val Ile Met Gly
 145 150 155 160
 Val Val Leu Gln Gly Ala Asn Leu Tyr Gly Tyr Ile Arg Cys Lys Val
 165 170 175
 Arg Ser Arg Lys His Leu Thr Ser Met Ala Thr Ser Tyr Phe Gly Lys
 180 185 190
 Gln Phe Leu Arg Gln Asn Thr Gly Asp Asp Gln Thr Ser
 195 200 205

<210> 64

<211> 213
 <212> PRT
 <213> Homo sapiens

<400> 64

Met Gly Asn Gly Thr Glu Glu Asp Tyr Asn Phe Val Phe Lys Val Val
 1 5 10 15
 Leu Ile Gly Glu Ser Gly Val Gly Lys Thr Asn Leu Leu Ser Arg Phe
 20 25 30
 Thr Arg Asn Glu Phe Ser His Asp Ser Arg Thr Thr Ile Gly Val Glu
 35 40 45
 Phe Ser Thr Arg Thr Val Met Leu Gly Thr Ala Ala Val Lys Ala Gln
 50 55 60
 Ile Trp Asp Thr Ala Gly Leu Glu Arg Tyr Arg Ala Ile Thr Ser Ala
 65 70 75 80
 Tyr Tyr Arg Gly Ala Val Gly Ala Leu Leu Val Phe Asp Leu Thr Lys
 85 90 95
 His Gln Thr Tyr Ala Val Val Glu Arg Trp Leu Lys Glu Leu Tyr Asp
 100 105 110
 His Ala Glu Ala Thr Ile Val Val Met Leu Val Gly Asn Lys Ser Asp
 115 120 125
 Leu Ser Gln Ala Arg Glu Val Pro Thr Glu Glu Ala Arg Met Phe Ala
 130 135 140
 Glu Asn Asn Gly Leu Leu Phe Leu Glu Thr Ser Ala Leu Asp Ser Thr
 145 150 155 160
 Asn Val Glu Leu Ala Phe Glu Thr Val Leu Lys Glu Ile Phe Ala Lys
 165 170 175
 Val Ser Lys Gln Arg Gln Asn Ser Ile Arg Thr Asn Ala Ile Thr Leu
 180 185 190
 Gly Ser Ala Gln Ala Gly Gln Glu Pro Gly Pro Gly Glu Lys Arg Ala
 195 200 205
 Cys Cys Ile Ser Leu
 210

<210> 65
 <211> 97
 <212> PRT
 <213> Homo sapiens

<400> 65

Met Leu Gly Asn Lys Arg Leu Gly Leu Ser Gly Leu Thr Leu Ala Leu
 1 5 10 15
 Ser Leu Leu Val Cys Leu Gly Ala Leu Ala Glu Ala Tyr Pro Ser Lys
 20 25 30
 Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp Met Ala Arg Tyr
 35 40 45
 Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr Arg Gln Arg Tyr
 50 55 60

Gly Lys Arg Ser Ser Pro Glu Thr Leu Ile Ser Asp Leu Leu Met Arg
65 70 75 80

Glu Ser Thr Glu Asn Val Pro Arg Thr Arg Leu Glu Asp Pro Ala Met
85 90 95

Trp

<210> 66

<211> 266

<212> PRT

<213> Homo sapiens

<400> 66

Met Gln Gly Ser Thr Arg Arg Met Gly Val Met Thr Asp Val His Arg
1 5 10 15

Arg Phe Leu Gln Leu Leu Met Thr His Gly Val Leu Glu Glu Trp Asp
20 25 30

Val Lys Arg Leu Gln Thr His Cys Tyr Lys Val His Asp Arg Asn Ala
35 40 45

Thr Val Asp Lys Leu Glu Asp Phe Ile Asn Asn Ile Asn Ser Val Leu
50 55 60

Glu Ser Leu Tyr Ile Glu Ile Lys Arg Gly Val Thr Glu Asp Asp Gly
65 70 75 80

Arg Pro Ile Tyr Ala Leu Val Asn Leu Ala Thr Thr Ser Ile Ser Lys
85 90 95

Met Ala Thr Asp Phe Ala Glu Asn Glu Leu Asp Leu Phe Arg Lys Ala
100 105 110

Leu Glu Leu Ile Ile Asp Ser Glu Thr Gly Phe Ala Ser Ser Thr Asn
115 120 125

Ile Leu Asn Leu Val Asp Gln Leu Lys Gly Lys Lys Met Arg Lys Lys
130 135 140

Glu Ala Glu Gln Val Leu Gln Lys Phe Val Gln Asn Lys Trp Leu Ile
145 150 155 160

Glu Lys Glu Gly Glu Phe Thr Leu His Gly Arg Ala Ile Leu Glu Met
165 170 175

Glu Gln Tyr Ile Arg Glu Thr Tyr Pro Asp Ala Val Lys Ile Cys Asn
180 185 190

Ile Cys His Ser Leu Leu Ile Gln Gly Gln Ser Cys Glu Thr Cys Gly
195 200 205

Ile Arg Met His Leu Pro Cys Val Ala Lys Tyr Phe Gln Ser Asn Ala
210 215 220

Glu Pro Arg Cys Pro His Cys Asn Asp Tyr Trp Pro His Glu Ile Pro
225 230 235 240

Lys Val Phe Asp Pro Glu Lys Glu Arg Glu Ser Gly Val Leu Lys Ser
245 250 255

Asn Lys Lys Ser Leu Arg Ser Arg Gln His
260 265

<210> 67
 <211> 149
 <212> PRT
 <213> Homo sapiens

<400> 67

```

Met Asn Tyr Leu Phe Phe Phe Leu Thr Thr Ser Gly Leu Tyr Cys Leu
 1              5              10              15

Ser Gly Ser His Gly Ser Asn Val Lys Tyr Ile Val Leu Thr Tyr Phe
              20              25              30

Asn Cys Ser Trp Ser Leu Thr Ser Pro Gly Phe Arg Asp Val Leu Lys
              35              40              45

Gly Ser Gln Leu Trp Gln Val Thr Asp Ser Trp Glu Met Glu Arg Thr
 50              55              60

Lys Glu Tyr Ser Ser Cys Leu Thr Phe Leu Pro Thr Ala Asp Ile Val
 65              70              75              80

Gln Ala Arg Val Met Glu Glu Leu Asn Leu Leu Ala Ser Gln Ala Ala
              85              90              95

Pro Ile Pro Thr Ser Gln Cys Thr Ala Pro Pro His Leu Phe Ser Pro
              100              105              110

Leu Ser Leu Thr Ser Pro Phe Ile Met Ser His Lys Ser Gly Thr Val
 115              120              125

Gly Ser His Tyr Asn Leu Leu Cys His Arg Asp Ser Ile Phe Leu Ile
 130              135              140

Ser Asn His Val Ser
145
  
```

<210> 68
 <211> 277
 <212> PRT
 <213> Homo sapiens

<400> 68

```

Met Ser Lys Lys Lys Ser Arg Gln Gly Lys Leu Val Lys Gln Phe Ala
 1              5              10              15

Lys Ile Glu Glu Ser Thr Pro Val His Asp Ser Pro Gly Lys Asp Asp
              20              25              30

Ala Val Pro Asp Leu Met Gly Pro His Ser Asp Gln Gly Glu His Ser
 35              40              45

Gly Thr Val Gly Val Pro Val Ser Tyr Thr Asp Cys Ala Pro Ser Pro
 50              55              60

Val Gly Cys Ser Val Val Thr Ser Asp Ser Phe Lys Thr Lys Asp Ser
 65              70              75              80

Phe Arg Thr Ala Lys Ser Lys Lys Lys Arg Arg Ile Thr Arg Tyr Asp
              85              90              95

Ala Gln Leu Ile Leu Glu Asn Asn Ser Gly Ile Pro Lys Leu Thr Leu
 100              105              110
  
```

Arg Arg Arg His Asp Ser Ser Ser Lys Thr Asn Asp Gln Glu Asn Asp
 115 120 125
 Gly Met Asn Ser Ser Lys Ile Ser Ile Lys Leu Ser Lys Asp His Asp
 130 135 140
 Asn Asp Asn Asn Leu Tyr Val Ala Lys Leu Asn Asn Gly Phe Asn Ser
 145 150 155 160
 Gly Ser Gly Ser Ser Ser Thr Lys Leu Lys Ile Gln Leu Lys Arg Asp
 165 170 175
 Glu Glu Asn Arg Gly Ser Tyr Thr Glu Gly Leu His Glu Asn Gly Val
 180 185 190
 Cys Cys Ser Asp Pro Leu Ser Leu Leu Glu Ser Arg Met Glu Val Asp
 195 200 205
 Asp Tyr Ser Gln Tyr Glu Glu Glu Ser Thr Asp Asp Ser Ser Ser Ser
 210 215 220
 Glu Gly Asp Glu Glu Glu Asp Asp Tyr Asp Asp Asp Phe Glu Asp Asp
 225 230 235 240
 Phe Ile Pro Leu Pro Pro Ala Lys Arg Leu Arg Leu Ile Val Gly Lys
 245 250 255
 Asp Ser Ile Asp Ile Asp Ile Ser Ser Arg Arg Arg Glu Asp Gln Ser
 260 265 270
 Leu Arg Leu Asn Ala
 275

<210> 69
 <211> 94
 <212> PRT
 <213> Homo sapiens

<400> 69
 Met His Ser Met Glu His Lys Leu Leu Trp Ile Leu Gln Leu Val Thr
 1 5 10 15
 Trp Asn Cys Phe Leu Val His Met Asn Thr Gly Ser Ile Gln Ala Gln
 20 25 30
 Leu Leu Pro Thr Ala Ser Leu Trp Ala Ser Cys Ser Gln Lys Ala Phe
 35 40 45
 His Leu Met Leu Pro Ile Ala Cys Leu Leu Ser Ser Arg Val Trp Pro
 50 55 60
 Ile Cys His Gly Glu Ala Ala Val Ser Lys Pro Ala Gly Asn Trp Asp
 65 70 75 80
 Val Ala Gly Asp Glu Arg Thr Asp Pro Ser Val Leu Pro Ala
 85 90

<210> 70
 <211> 449
 <212> PRT
 <213> Homo sapiens

<400> 70
 Met Gln Phe Ala Trp Gln Ser Tyr Lys Arg Tyr Ala Met Gly Lys Asn

1	5	10	15
Glu Leu Arg Pro 20	Leu Thr Lys Asp Gly 25	Tyr Glu Gly Asn Met 30	Phe Gly
Gly Leu Ser 35	Gly Ala Thr Val Ile Asp 40	Ser Leu Asp Thr 45	Leu Tyr Leu
Met Glu Leu Lys 50	Glu Glu Phe 55	Gln Glu Ala Lys 60	Trp Val Gly Glu
Ser Phe His Leu 65	Asn Val Ser Gly 70	Glu Ala Ser Leu 75	Phe Glu Val Asn 80
Ile Arg Tyr Ile 85	Gly Gly Leu Leu Ser 90	Ala Phe Tyr Leu Thr 95	Gly Glu
Glu Val Phe Arg 100	Ile Lys Ala Ile Arg 105	Leu Gly Glu Lys 110	Leu Leu Pro
Ala Phe Asn Thr 115	Pro Thr Gly Ile 120	Pro Lys Gly Val 125	Val Ser Phe Lys
Ser Gly Asn Trp 130	Gly Trp Ala Thr 135	Ala Gly Ser Ser 140	Ser Ile Leu Ala
Glu Phe Gly Ser 145	Leu His Leu Glu Phe 150	Leu His Leu Thr 155	Glu Leu Ser 160
Gly Asn Gln Val 165	Phe Ala Glu Lys Val 170	Arg Asn Ile Arg 175	Lys Val Leu
Arg Lys Ile 180	Glu Lys Pro Phe Gly 185	Leu Tyr Pro Asn Phe 190	Leu Ser Pro
Val Ser Gly 195	Asn Trp Val Gln His 200	His Val Ser Val 205	Gly Gly Leu Gly
Asp Ser Phe Tyr 210	Glu Tyr Leu Ile Lys 215	Ser Trp Leu Met 220	Ser Gly Lys
Thr Asp Met Glu 225	Ala Lys Asn Met Tyr 230	Tyr Tyr Glu Ala 235	Leu Glu Ala Ile 240
Glu Thr Tyr Leu 245	Leu Asn Val Ser Pro Gly 250	Gly Leu Thr Tyr 255	Ile Ala
Glu Trp Arg Gly 260	Gly Ile Leu Asp His 265	Lys Met Gly His 270	Leu Ala Cys
Phe Ser Gly Gly 275	Met Ile Ala Leu Gly 280	Ala Glu Asp Ala 285	Lys Glu Glu
Lys Arg Ala His 290	Tyr Arg Glu Leu Ala 295	Ala Gln Ile Thr 300	Lys Thr Cys
His Glu Ser Tyr 305	Ala Arg Ser Asp Thr 310	Lys Leu Gly Pro 315	Glu Ala Phe 320
Trp Phe Asn Ser 325	Gly Arg Glu Ala Val 330	Ala Thr Gln Leu Ser 335	Glu Ser
Tyr Tyr Ile 340	Leu Arg Pro Glu Val Val 345	Glu Ser Tyr Met Tyr 350	Leu Trp
Arg Gln Thr His Asn Pro Ile Tyr Arg Glu Trp Gly Trp Glu Val Val			

355 360 365
 Leu Ala Leu Glu Lys Tyr Cys Arg Thr Glu Ala Gly Phe Ser Gly Ile
 370 375 380
 Gln Asp Val Tyr Ser Ser Thr Pro Asn His Asp Asn Lys Gln Gln Ser
 385 390 395 400
 Phe Phe Leu Ala Glu Thr Leu Lys Tyr Leu Tyr Leu Leu Phe Ser Glu
 405 410 415
 Asp Asp Leu Leu Ser Leu Glu Asp Trp Val Phe Asn Thr Glu Ala His
 420 425 430
 Pro Leu Pro Val Asn His Ser Asp Ser Ser Gly Arg Ala Trp Gly Arg
 435 440 445
 His

<210> 71
 <211> 372
 <212> PRT
 <213> Homo sapiens

<400> 71
 Met Thr Phe Gln Phe Asn Phe Thr Ile Glu Asp His Leu Glu Asn Glu
 1 5 10 15
 Leu Thr Pro Ile Arg Asp Gly Ala Leu Thr Leu Asp Ser Ser Lys Glu
 20 25 30
 Leu Ser Val Ser Glu Ser Gln Lys Gly Glu Glu Arg Asp Arg Lys Cys
 35 40 45
 Ser Ala Glu Gln Phe Asp Leu Pro Gln Asp His Leu Trp Glu His Lys
 50 55 60
 Ser Met Glu Asn Ala Ala Pro Ser Gln Asp Thr Asp Ser Pro Leu Ser
 65 70 75 80
 Ala Ala Ser Ser Ser Arg Asn Leu Glu Pro His Gly Lys Gln Pro Ser
 85 90 95
 Leu Arg Ala Ala Lys Glu His Ala Met Pro Lys Asp Leu Lys Lys Met
 100 105 110
 Leu Glu Asn Lys Val Ile Glu Thr Leu Pro Gly Phe Gln His Val Lys
 115 120 125
 Leu Ser Val Val Lys Thr Ile Leu Leu Lys Glu Asn Phe Pro Gly Glu
 130 135 140
 Asn Ile Val Ser Lys Ser Phe Ser Ser His Ser Asp Leu Ile Thr Gly
 145 150 155 160
 Val Tyr Glu Gly Gly Leu Lys Ile Trp Glu Cys Thr Phe Asp Leu Leu
 165 170 175
 Ala Tyr Phe Thr Lys Ala Lys Val Lys Phe Ala Gly Lys Lys Val Leu
 180 185 190
 Asp Leu Gly Cys Gly Ser Gly Leu Leu Gly Ile Thr Ala Phe Lys Gly
 195 200 205

Gly Ser Lys Glu Ile His Phe Gln Asp Tyr Asn Ser Met Val Ile Asp
 210 215 220
 Glu Val Thr Leu Pro Asn Val Val Ala Asn Ser Thr Leu Glu Asp Glu
 225 230 235 240
 Glu Asn Asp Val Asn Glu Pro Asp Val Lys Arg Cys Arg Lys Pro Lys
 245 250 255
 Val Thr Gln Leu Tyr Lys Cys Arg Phe Phe Ser Gly Glu Trp Ser Glu
 260 265 270
 Phe Cys Lys Leu Val Leu Ser Ser Glu Lys Leu Phe Val Lys Tyr Asp
 275 280 285
 Leu Ile Leu Thr Ser Glu Thr Ile Tyr Asn Pro Asp Tyr Tyr Ser Asn
 290 295 300
 Leu His Gln Thr Phe Leu Arg Leu Leu Ser Lys Asn Gly Arg Val Leu
 305 310 315 320
 Leu Ala Ser Lys Ala His Tyr Phe Gly Val Gly Gly Gly Val His Leu
 325 330 335
 Phe Gln Lys Phe Val Glu Glu Arg Asp Val Phe Lys Thr Arg Ile Leu
 340 345 350
 Lys Ile Ile Asp Glu Gly Leu Lys Arg Phe Ile Ile Glu Ile Thr Phe
 355 360 365
 Lys Phe Pro Gly
 370

<210> 72
 <211> 211
 <212> PRT
 <213> Homo sapiens

<400> 72
 Met Thr Thr Leu Thr Arg Gln Asp Leu Asn Phe Gly Gln Val Val Ala
 1 5 10 15
 Asp Val Leu Cys Glu Phe Leu Glu Val Ala Val His Leu Ile Leu Tyr
 20 25 30
 Val Arg Glu Val Tyr Pro Val Gly Ile Phe Gln Lys Arg Lys Lys Tyr
 35 40 45
 Asn Val Pro Val Gln Met Ser Cys His Pro Glu Leu Asn Gln Tyr Ile
 50 55 60
 Gln Asp Thr Leu His Cys Val Lys Pro Leu Leu Glu Lys Asn Asp Val
 65 70 75 80
 Glu Lys Val Val Val Ile Leu Asp Lys Glu His Arg Pro Val Glu
 85 90 95
 Lys Phe Val Phe Glu Ile Thr Gln Pro Pro Leu Leu Ser Ile Ser Ser
 100 105 110
 Asp Ser Leu Leu Ser His Val Glu Gln Leu Leu Arg Ala Phe Ile Leu
 115 120 125
 Lys Ile Ser Val Cys Asp Ala Val Leu Asp His Asn Pro Pro Gly Cys
 130 135 140

Thr Phe Thr Val Leu Val His Thr Arg Glu Ala Ala Thr Arg Asn Met
145 150 155 160

Glu Lys Ile Gln Val Ile Lys Asp Phe Pro Trp Ile Leu Ala Asp Glu
165 170 175

Gln Asp Val His Met His Asp Pro Arg Leu Ile Pro Leu Lys Thr Met
180 185 190

Thr Ser Asp Ile Leu Lys Met Gln Leu Tyr Val Glu Glu Arg Ala His
195 200 205

Lys Gly Ser
210

<210> 73

<211> 219

<212> PRT

<213> Homo sapiens

<400> 73

Ala Val Val Gly Tyr Thr Asn Cys Gly Lys Thr Thr Leu Ile Lys Ala
1 5 10 15

Leu Thr Gly Asp Ala Ala Ile Gln Pro Arg Asp Gln Leu Phe Ala Thr
20 25 30

Leu Asp Val Thr Ala His Ala Gly Thr Leu Pro Ser Arg Met Thr Val
35 40 45

Leu Tyr Val Asp Thr Ile Gly Phe Leu Ser Gln Leu Pro His Gly Leu
50 55 60

Ile Glu Ser Phe Ser Ala Thr Leu Glu Asp Val Ala His Ser Asp Leu
65 70 75 80

Ile Leu His Val Arg Asp Val Ser His Pro Glu Ala Glu Leu Gln Lys
85 90 95

Cys Ser Val Leu Ser Thr Leu Arg Gly Leu Gln Leu Pro Ala Pro Leu
100 105 110

Leu Asp Ser Met Val Glu Val His Asn Lys Val Asp Leu Val Pro Gly
115 120 125

Tyr Ser Pro Thr Glu Pro Asn Val Val Pro Val Ser Ala Leu Arg Gly
130 135 140

His Gly Leu Gln Glu Leu Lys Ala Glu Leu Asp Ala Ala Val Leu Lys
145 150 155 160

Ala Thr Gly Arg Gln Ile Leu Thr Leu Arg Val Arg Leu Ala Gly Ala
165 170 175

Gln Leu Ser Trp Leu Tyr Lys Glu Ala Thr Val Gln Glu Val Asp Val
180 185 190

Ile Pro Glu Asp Gly Ala Ala Asp Val Arg Val Ile Ile Ser Asn Ser
195 200 205

Ala Tyr Gly Lys Phe Arg Lys Leu Phe Pro Gly
210 215

<210> 74
 <211> 221
 <212> PRT
 <213> Homo sapiens

<400> 74
 Met Leu Ser Leu Pro Leu Arg Ala Pro Ala Pro Arg Leu Glu Arg Arg
 1 5 10 15
 Pro Ala Gly Pro Pro Ala Asp Val Phe Leu Val Pro Lys Arg Val Val
 20 25 30
 Arg Ala Ser Arg Pro Leu Arg Asp Leu Arg Ala Ser His Arg Ala Pro
 35 40 45
 Arg Thr Gln Arg Ala Trp Ser Ser Pro Leu Thr Pro Ser Pro Ala Gly
 50 55 60
 Thr His Ala Gly Ser Thr His Thr Ser Ala Pro Pro Pro Asn Phe Trp
 65 70 75 80
 Glu Arg Thr Pro Gly Ser Ala Gln Pro Leu Ala Phe Gln Lys Pro Leu
 85 90 95
 Tyr Ala Tyr Leu Ile Phe Val Ile Gly Asp Glu Pro Ser Leu Leu Ser
 100 105 110
 Pro Phe Pro His Thr His Gln Ser Pro Leu Ala Ile Pro Ser Pro Ser
 115 120 125
 Ala Ser Pro Pro Pro Ser Cys Ala Pro Ala Pro His Ser His Pro Pro
 130 135 140
 Pro Ile Gly Leu Ala Leu Ala Cys Lys Ser Arg Arg Trp Pro Arg Ala
 145 150 155 160
 Gln Pro Ser Arg Met Ser Pro Gly Pro Pro Leu Trp Glu Arg Arg Gln
 165 170 175
 Ser Tyr Trp Pro Leu Thr Arg Pro Leu Gly Pro Arg Ala Arg Gln Ala
 180 185 190
 Phe Glu Ser Thr Cys Ser Ser Pro Glu Ser Arg Pro Arg Pro Cys Leu
 195 200 205
 Pro His Arg Ser Arg Pro Gln Ser Thr Leu Pro Gln Leu
 210 215 220

<210> 75
 <211> 48
 <212> PRT
 <213> Homo sapiens

<400> 75
 Met Leu Cys Trp Phe Cys Phe Phe Val Leu Leu Phe Phe Phe Phe
 1 5 10 15
 Leu Phe Cys Phe Leu Val Met Trp Leu Lys Cys Asn Ser Phe Phe Phe
 20 25 30
 Gly Thr Tyr Phe Cys Gln Leu Lys Thr Arg Arg Ala Gln Leu Phe Phe
 35 40 45

<210> 76
 <211> 63
 <212> PRT
 <213> Homo sapiens

<400> 76
 Met Lys Thr Gly Gly Lys His Ser Val Ile Arg Tyr Phe Ser Asn Ile
 1 5 10 15
 Lys Thr Thr Lys Thr Asn Asp Lys Asn Val Tyr Phe Tyr Thr Pro Ala
 20 25 30
 Tyr Arg Val Ser Phe Arg Asp Val Tyr Glu Tyr Leu Asn Leu Leu Ile
 35 40 45
 Ser Val Leu Met Lys Ala Glu Leu Asn Arg Lys Ala Ser Ser Trp
 50 55 60

<210> 77
 <211> 36
 <212> PRT
 <213> Homo sapiens

<400> 77
 Met Phe Ser Val Leu Cys Leu Cys Val Cys Ala Arg Gln Arg Asp Arg
 1 5 10 15
 Leu Phe Val Lys Ser Ala Ser Cys Leu Gly Ile Phe Val Ser His Leu
 20 25 30
 Ala Val Ser Ser
 35

<210> 78
 <211> 395
 <212> PRT
 <213> Homo sapiens

<400> 78
 Met Val Ser Ser Ser Asn Leu Pro Gln Gly Trp Leu Glu Val Gln Gly
 1 5 10 15
 Ile Pro Glu Gly Trp Asp Gly Val Ala Gly Trp Tyr Leu Pro Gly Ile
 20 25 30
 Asn Pro Gly Arg Thr Ala Arg Arg Phe Ala Tyr Leu Phe Val Asn Ile
 35 40 45
 Asn Val Thr Ser Glu Pro His Glu Val Leu Ala Leu Trp Phe Leu Trp
 50 55 60
 Tyr Val Lys Gln Cys Gly Gly Thr Thr Arg Ile Phe Ser Val Thr Asn
 65 70 75 80
 Gly Gly Gln Glu Arg Lys Phe Val Gly Gly Ser Gly Gln Val Ser Glu
 85 90 95
 Arg Ile Met Asp Leu Leu Gly Asp Gln Val Lys Leu Asn His Pro Val
 100 105 110
 Thr His Val Asp Gln Ser Ser Asp Asn Ile Ile Ile Glu Thr Leu Asn
 115 120 125

His Glu His Tyr Glu Cys Lys Tyr Val Ile Asn Ala Ile Pro Pro Thr
 130 135 140
 Leu Thr Ala Lys Ile His Phe Arg Pro Glu Leu Pro Ala Glu Arg Asn
 145 150 155 160
 Gln Leu Ile Gln Arg Leu Pro Met Gly Ala Val Ile Lys Cys Met Met
 165 170 175
 Tyr Tyr Lys Glu Ala Phe Trp Lys Lys Lys Asp Tyr Cys Gly Cys Met
 180 185 190
 Ile Ile Glu Asp Glu Asp Ala Pro Ile Ser Ile Thr Leu Asp Asp Thr
 195 200 205
 Lys Pro Asp Gly Ser Leu Pro Ala Ile Met Gly Phe Ile Leu Ala Arg
 210 215 220
 Lys Ala Asp Arg Leu Ala Lys Leu His Lys Glu Ile Arg Lys Lys Lys
 225 230 235 240
 Ile Cys Glu Leu Tyr Ala Lys Val Leu Gly Ser Gln Glu Ala Leu His
 245 250 255
 Pro Val His Tyr Glu Glu Lys Asn Trp Cys Glu Glu Gln Tyr Ser Gly
 260 265 270
 Gly Cys Tyr Thr Ala Tyr Phe Pro Pro Gly Ile Met Thr Gln Tyr Gly
 275 280 285
 Arg Val Ile Arg Gln Pro Val Gly Arg Ile Phe Phe Ala Gly Thr Glu
 290 295 300
 Thr Ala Thr Lys Trp Ser Gly Tyr Met Glu Gly Ala Val Glu Ala Gly
 305 310 315 320
 Glu Arg Ala Ala Arg Glu Val Leu Asn Gly Leu Gly Lys Val Thr Glu
 325 330 335
 Lys Asp Ile Trp Val Gln Glu Pro Glu Ser Lys Asp Val Pro Ala Val
 340 345 350
 Glu Ile Thr His Thr Phe Trp Glu Arg Asn Leu Pro Ser Val Ser Gly
 355 360 365
 Leu Leu Lys Ile Ile Gly Phe Ser Thr Ser Val Thr Ala Leu Gly Phe
 370 375 380
 Val Leu Tyr Lys Tyr Lys Leu Leu Pro Arg Ser
 385 390 395

<210> 79

<211> 45

<212> PRT

<213> Homo sapiens

<400> 79

Met Gly Asn Cys Ser Leu Leu Leu Pro Thr Leu Ser Glu His Leu Trp
 1 5 10 15
 Asp Leu Val Ala Gln Asn Ser Ser Val Thr Ser Ser Ser Pro Pro Phe
 20 25 30
 Val Pro Ile Ser Ser Leu Ser Lys Lys Pro Cys Gly Thr

45

35

40

45

<210> 80
 <211> 79
 <212> PRT
 <213> Homo sapiens

<400> 80
 Met Trp Gly Ser Cys Val Leu Glu Tyr Tyr Val Ser Pro Pro Ser Ala
 1 5 10 15
 Val Phe Ser Glu His Val Cys Cys Pro Trp Trp Glu Arg Gly His Cys
 20 25 30
 Ala Val Val His Arg Cys Leu Ser Phe Thr Val Gly Leu Ser Val Cys
 35 40 45
 Leu Ser Phe Leu Ser Ala Ala Gln Met Glu Asn Asn Tyr Leu Leu His
 50 55 60
 Trp Arg Glu Arg Lys Ser Leu Arg Ile Pro Lys Gly Thr Leu Ala
 65 70 75

<210> 81
 <211> 36
 <212> PRT
 <213> Homo sapiens

<400> 81
 Met Cys Pro Phe Ser Ser Leu His Leu Ala Ala Gly Ile Val Asp Ile
 1 5 10 15
 Thr Gly Ala Leu Ala Ala Val Ser Arg Gly Ser Lys Pro His Pro Lys
 20 25 30
 Ser Lys Ala Asp
 35

<210> 82
 <211> 35
 <212> PRT
 <213> Homo sapiens

<400> 82
 Met Ala Leu Gly Thr Trp Lys Arg Val Thr Glu Met Gly Arg His Glu
 1 5 10 15
 Leu Lys Glu Ala Ile Tyr Asp His Val Ile Cys Asn Met Lys Lys Ala
 20 25 30
 Tyr Leu Glu
 35

<210> 83
 <211> 144
 <212> PRT
 <213> Homo sapiens

<400> 83
 Met Glu Ala Trp Ile Arg Ala Asn Gln Pro Ala Phe Leu Val Trp Arg
 1 5 10 15

Ser Thr Trp Pro Phe Pro Trp Ala Gln Gly His Leu Lys His Cys Pro
 20 25 30
 Val Lys Leu Val Leu Gly Cys Pro Cys Ala Trp Arg Val Leu Lys Leu
 35 40 45
 Thr Phe Gln Ile Pro Arg Glu Gln Gly Glu Ile Ser Arg Met Ser Ile
 50 55 60
 Ala Ala Lys Lys Cys Leu Gly Gly Leu Pro Leu Leu Thr Pro His Leu
 65 70 75 80
 Ala Ala Asp Gln His Ser Ile Leu Asn Thr Leu Arg Ala Pro Ser Met
 85 90 95
 Ala Phe Asp Arg Thr Lys Ser Pro Gly Val Val Thr Glu Asn Arg Ser
 100 105 110
 Cys Ala Val Thr Ala Met Phe Pro Pro Gly Arg Gln Lys Leu Lys Ser
 115 120 125
 Pro Lys Arg Thr Ser Phe Ser Ser Ala Ala Asp Glu Trp His Arg Tyr
 130 135 140

<210> 84
 <211> 369
 <212> PRT
 <213> Homo sapiens

<400> 84
 Met Pro Arg Ala Pro Lys Arg Gln Arg Cys Met Pro Glu Glu Asp Leu
 1 5 10 15
 Gln Ser Gln Ser Glu Thr Gln Gly Leu Glu Gly Ala Gln Ala Pro Leu
 20 25 30
 Ala Val Glu Glu Asp Ala Ser Ser Ser Thr Ser Thr Ser Ser Ser Phe
 35 40 45
 Pro Ser Ser Phe Pro Ser Ser Ser Ser Ser Ser Ser Ser Ser Cys Tyr
 50 55 60
 Pro Leu Ile Pro Ser Thr Pro Glu Glu Val Ser Ala Asp Asp Glu Thr
 65 70 75 80
 Pro Asn Pro Pro Gln Ser Ala Gln Ile Ala Cys Ser Ser Pro Ser Val
 85 90 95
 Val Ala Ser Leu Pro Leu Asp Gln Ser Asp Glu Gly Ser Ser Ser Gln
 100 105 110
 Lys Glu Glu Ser Pro Ser Thr Leu Gln Val Leu Pro Asp Ser Glu Ser
 115 120 125
 Leu Pro Arg Ser Glu Ile Asp Glu Lys Val Thr Asp Leu Val Gln Phe
 130 135 140
 Leu Leu Phe Lys Tyr Gln Met Lys Glu Pro Ile Thr Lys Ala Glu Ile
 145 150 155 160
 Leu Glu Ser Val Ile Lys Asn Tyr Glu Asp His Phe Pro Leu Leu Phe
 165 170 175

Ser Glu Ala Ser Glu Cys Met Leu Leu Val Phe Gly Ile Asp Val Lys
 180 185 190
 Glu Val Asp Pro Thr Gly His Ser Phe Val Leu Val Thr Ser Leu Gly
 195 200 205
 Leu Thr Tyr Asp Gly Met Leu Ser Asp Val Gln Ser Met Pro Lys Thr
 210 215 220
 Gly Ile Leu Ile Leu Ile Leu Ser Ile Ile Phe Ile Glu Gly Tyr Cys
 225 230 235 240
 Thr Pro Glu Glu Val Ile Trp Glu Ala Leu Asn Met Met Gly Leu Tyr
 245 250 255
 Asp Gly Met Glu His Leu Ile Tyr Gly Glu Pro Arg Lys Leu Leu Thr
 260 265 270
 Gln Asp Trp Val Gln Glu Asn Tyr Leu Glu Tyr Arg Gln Val Pro Gly
 275 280 285
 Ser Asp Pro Ala Arg Tyr Glu Phe Leu Trp Gly Pro Arg Ala His Ala
 290 295 300
 Glu Ile Arg Lys Met Ser Leu Leu Lys Phe Leu Ala Lys Val Asn Gly
 305 310 315 320
 Ser Asp Pro Arg Ser Phe Pro Leu Trp Tyr Glu Glu Ala Leu Lys Asp
 325 330 335
 Glu Glu Glu Arg Ala Gln Asp Arg Ile Ala Thr Thr Asp Asp Thr Thr
 340 345 350
 Ala Met Ala Ser Ala Ser Ser Ser Ala Thr Gly Ser Phe Ser Tyr Pro
 355 360 365
 Glu

<210> 85

<211> 69

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (8)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 85

Leu Val Ile Tyr Ser Trp His Xaa Phe Phe Ser Phe Gly Phe Ala Trp
 1 5 10 15

Leu Phe Leu Gln Val Leu Ser Arg Tyr His Ser Ala Asn His Cys Tyr
 20 25 30

Arg Met Val Thr Ser Phe Val Leu Thr Val Gln Gln Gln Ile Trp Val
 35 40 45

Arg Leu Asn Leu Ser Val Asn Phe Phe Phe Trp Cys Phe Phe Gly Leu
 50 55 60

Met Thr Val Ser Leu
 65

<210> 86
 <211> 95
 <212> PRT
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (46)
 <223> Xaa equals any of the naturally occurring L-amino acids

<400> 86
 Pro Val Leu Leu Ser Leu Leu Leu Leu Leu Gly Pro Ala Val Pro Gln
 1 5 10 15
 Glu Asn Gln Asp Gly Arg Tyr Ser Leu Thr Tyr Ile Tyr Thr Gly Leu
 20 25 30
 Ser Lys His Val Glu Asp Val Pro Ala Phe Gln Ala Leu Xaa His Ser
 35 40 45
 Met Thr Ser Ser Ser Leu Asp Thr Thr Val Lys Thr Gly Ser Leu Ser
 50 55 60
 Pro Trp Asp Ser Gly Asp Arg Trp Lys Glu Trp Arg Ile Gly Ser Arg
 65 70 75 80
 Thr Ala Asn Phe Arg Arg Pro Gly Arg Thr Ser Leu Trp Arg Pro
 85 90 95

<210> 87
 <211> 181
 <212> PRT
 <213> Homo sapiens

<400> 87
 Met Leu Gln Gln Asp Ser Asn Asp Asp Thr Glu Asp Val Ser Leu Phe
 1 5 10 15
 Asp Ala Glu Glu Glu Thr Thr Asn Arg Pro Arg Lys Ala Lys Ile Arg
 20 25 30
 His Pro Val Ala Ser Phe Phe His Leu Phe Phe Arg Val Ser Ala Ile
 35 40 45
 Ile Val Tyr Leu Leu Cys Glu Leu Leu Ser Ser Ser Phe Ile Thr Cys
 50 55 60
 Met Val Thr Ile Ile Leu Leu Leu Ser Cys Asp Phe Trp Ala Val Lys
 65 70 75 80
 Asn Val Thr Gly Arg Leu Met Val Gly Leu Arg Trp Trp Asn His Ile
 85 90 95
 Asp Glu Asp Gly Lys Ser His Trp Val Phe Glu Ser Arg Lys Glu Ser
 100 105 110
 Ser Gln Glu Asn Lys Thr Val Ser Glu Ala Glu Ser Arg Ile Phe Trp
 115 120 125
 Leu Gly Leu Ile Ala Cys Pro Val Leu Trp Val Ile Phe Ala Phe Ser
 130 135 140
 Ala Leu Phe Ser Phe Arg Val Lys Trp Leu Ala Val Val Ile Met Gly

145		150		155		160
Val Val Leu Gln Gly Ala Asn Leu Tyr Gly Thr Ser Gly Val Arg Cys						
		165		170		175
Ala Ala Glu Ser Ile						
		180				

```
<210> 88
<211> 61
<212> PRT
<213> Homo sapiens
```

```
<220>
<221> SITE
<222> (39)
<223> Xaa equals any of the naturally occurring L-amino acids
```

```
<220>
<221> SITE
<222> (54)
<223> Xaa equals any of the naturally occurring L-amino acids
```

```

<400> 88
Leu Cys Leu Gln Gly Tyr Tyr Arg Gly Ala Val Gly Ala Leu Leu Val
  1             5             10             15
Phe Asp Leu Thr Lys His Gln Thr Tyr Ala Val Val Glu Arg Trp Leu
      20             25             30
Lys Glu Leu Tyr Asp His Xaa Glu Ala Thr Ile Val Val Met Leu Val
      35             40             45
Gly Asn Lys Met Thr Xaa Ala Arg Pro Gly Lys Cys Pro
      50             55             60

```

```
<210> 89
<211> 217
<212> PRT
<213> Homo sapiens
```

```

<220>
<221> SITE
<222> (3)
<223> Xaa equals any of the naturally occurring L-amino acids

```

<400>	89															
Met	Arg	Xaa	Lys	Met	Gly	Asn	Gly	Thr	Glu	Glu	Asp	Tyr	Asn	Phe	Val	
1				5					10					15		
Phe	Lys	Val	Val	Leu	Ile	Gly	Glu	Ser	Gly	Val	Gly	Lys	Thr	Asn	Leu	
			20					25					30			
Leu	Ser	Arg	Phe	Thr	Arg	Asn	Glu	Phe	Ser	His	Asp	Ser	Arg	Thr	Thr	
		35					40					45				
Ile	Gly	Val	Glu	Phe	Ser	Thr	Arg	Thr	Val	Met	Leu	Gly	Thr	Ala	Ala	
	50					55					60					
Val	Lys	Ala	Gln	Ile	Trp	Asp	Thr	Ala	Gly	Leu	Glu	Arg	Tyr	Arg	Ala	
65					70					75					80	
Ile	Thr	Ser	Ala	Tyr	Tyr	Arg	Gly	Ala	Val	Gly	Ala	Leu	Leu	Val	Phe	
				85					90					95		

Asp Leu Thr Lys His Gln Thr Tyr Ala Val Val Glu Arg Trp Leu Lys
 100 105 110
 Glu Leu Tyr Asp His Ala Glu Ala Thr Ile Val Val Met Leu Val Gly
 115 120 125
 Asn Lys Ser Asp Leu Ser Gln Ala Arg Glu Val Pro Thr Glu Glu Ala
 130 135 140
 Arg Met Phe Ala Glu Asn Asn Gly Leu Leu Phe Leu Glu Thr Ser Ala
 145 150 155 160
 Leu Asp Ser Thr Asn Val Glu Leu Ala Phe Glu Thr Val Leu Lys Glu
 165 170 175
 Ile Phe Ala Lys Val Ser Lys Gln Arg Gln Asn Ser Ile Arg Thr Asn
 180 185 190
 Ala Ile Thr Ser Gly Ser Ala Gln Ala Gly Gln Glu Pro Gly Pro Gly
 195 200 205
 Glu Lys Arg Ala Cys Cys Ile Ser Leu
 210 215

<210> 90
 <211> 72
 <212> PRT
 <213> Homo sapiens

<400> 90
 Met Leu Gly Asn Lys Arg Leu Gly Leu Ser Gly Leu Thr Ser Pro Cys
 1 5 10 15
 Pro Cys Ser Cys Ala Trp Val Arg Trp Pro Arg Arg Thr Pro Pro Ser
 20 25 30
 Arg Thr Thr Arg Ala Arg Thr His Gln Arg Arg Thr Trp Pro Asp Thr
 35 40 45
 Thr Gln Arg Cys Asp Thr Thr Ser Thr Ser Ser Pro Gly Arg Asp Met
 50 55 60
 Glu Asn Asp Leu Ala Gln Arg His
 65 70

<210> 91
 <211> 91
 <212> PRT
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (32)
 <223> Xaa equals any of the naturally occurring L-amino acids

<220>
 <221> SITE
 <222> (42)
 <223> Xaa equals any of the naturally occurring L-amino acids

<220>
 <221> SITE
 <222> (51)

<223> Xaa equals any of the naturally occurring L-amino acids

<220>

<221> SITE

<222> (69)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 91

Met Leu Gly Asn Lys Arg Leu Gly Leu Ser Gly Leu Thr Leu Ala Leu
1 5 10 15

Ser Leu Leu Val Cys Leu Gly Ala Leu Ala Glu Ala Tyr Pro Ser Xaa
20 25 30

Pro Asp Asn Pro Gly Glu Asp Ala Pro Xaa Glu Gly His Gly Gln Ile
35 40 45

Leu Leu Xaa Ala Ala Thr Leu His Gln Pro His His Gln Ala Glu Ile
50 55 60

Trp Lys Thr Ile Xaa Pro Arg Asp Thr Asp Phe Arg Pro Leu Asp Glu
65 70 75 80

Arg Lys His Arg Lys Cys Ser Gln Asn Ser Ala
85 90

<210> 92

<211> 277

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (147)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 92

Met Gln Gly Ser Thr Arg Arg Met Gly Val Met Thr Asp Val His Arg
1 5 10 15

Arg Phe Leu Gln Leu Leu Met Thr His Gly Val Leu Glu Glu Trp Asp
20 25 30

Val Lys Arg Leu Gln Thr His Cys Tyr Lys Val His Asp Arg Asn Ala
35 40 45

Thr Val Asp Lys Leu Glu Asp Phe Ile Asn Asn Ile Asn Ser Val Leu
50 55 60

Glu Ser Leu Tyr Ile Glu Ile Lys Arg Gly Val Thr Glu Asp Asp Gly
65 70 75 80

Arg Pro Ile Tyr Ala Leu Val Asn Leu Ala Thr Thr Ser Ile Ser Lys
85 90 95

Met Ala Thr Asp Phe Ala Glu Asn Glu Leu Asp Leu Phe Arg Lys Ala
100 105 110

Leu Glu Leu Ile Ile Asp Ser Glu Thr Gly Phe Ala Ser Ser Thr Asn
115 120 125

Ile Leu Asn Leu Val Asp Gln Leu Lys Gly Lys Lys Met Arg Lys Lys
130 135 140

Glu Ala Xaa Gln Val Leu Gln Lys Phe Val Gln Asn Lys Trp Leu Ile

145		150		155		160
Glu Lys Glu Gly	Glu Phe Thr Leu His Gly	Arg Ala Ile Leu Glu Met				
	165	170			175	
Glu Gln Tyr Ile	Arg Glu Thr Tyr Pro Asp Ala Val Lys Ile Cys Asn					
	180	185			190	
Ile Cys His Ser	Leu Leu Ile Gln Gly Gln Ser Cys Glu Thr Cys Gly					
	195	200			205	
Ile Arg Met His	Leu Pro Cys Val Ala Lys Tyr Phe Gln Ser Asn Ala					
	210	215			220	
Glu Pro Arg Cys	Pro His Cys Asn Asp Tyr Trp Pro His Glu Ile Pro					
	225	230			235	240
Lys Val Phe Asp	Pro Glu Lys Glu Arg Glu Ser Gly Val Leu Lys Ser					
	245	250			255	
Asn Lys Lys Ser	Cys Gly Pro Gly Ser Ile Ser His Arg Ala Leu Leu					
	260	265			270	
Arg Gly Trp Leu Pro						
	275					

<210> 93

<211> 122

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (16)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 93

Phe Leu His Thr	Phe Asn Cys Ser Trp Ser Leu Thr Ser Pro Gly Xaa
1	5 10 15

Arg Asp Val Leu Lys Gly Ser Gln Leu Trp Gln Val Thr Asp Ser Trp
20 25 30

Glu Met Glu Arg Thr Lys Glu Tyr Ser Ser Cys Leu Thr Phe Leu Pro
35 40 45

Thr Ala Asp Ile Val Gln Ala Arg Val Met Glu Glu Leu Asn Leu Leu
50 55 60

Ala Ser Gln Ala Ala Pro Ile Pro Thr Ser Gln Cys Thr Ala Pro Pro
65 70 75 80

His Leu Phe Ser Pro Leu Ser Leu Thr Ser Pro Phe Ile Met Ser His
85 90 95

Lys Ser Gly Thr Val Gly Ser His Tyr Asn Leu Leu Cys His Arg Asp
100 105 110

Ser Ile Phe Leu Ile Ser Asn His Val Ser
115 120

<210> 94

<211> 341

<212> PRT

<213> Homo sapiens

<400> 94

Arg Thr Asn Leu Lys Glu Ala Ser Asp Ile Lys Leu Glu Pro Asn Thr
 1 5 10 15
 Leu Asn Gly Tyr Lys Ser Ser Val Thr Glu Pro Cys Pro Asp Ser Gly
 20 25 30
 Glu Gln Leu Gln Pro Ala Pro Val Leu Gln Glu Glu Leu Ala His
 35 40 45
 Glu Thr Ala Gln Lys Gly Glu Ala Lys Cys His Lys Ser Asp Thr Gly
 50 55 60
 Met Ser Lys Lys Lys Ser Arg Gln Gly Lys Leu Val Lys Gln Phe Ala
 65 70 75 80
 Lys Ile Glu Glu Ser Thr Pro Val His Asp Ser Pro Gly Lys Asp Asp
 85 90 95
 Ala Val Pro Asp Leu Met Gly Pro His Ser Asp Gln Gly Glu His Ser
 100 105 110
 Gly Thr Val Gly Val Pro Val Ser Tyr Thr Asp Cys Ala Pro Ser Pro
 115 120 125
 Val Gly Cys Ser Val Val Thr Ser Asp Ser Phe Arg Thr Lys Asp Ser
 130 135 140
 Phe Arg Thr Ala Lys Ser Lys Lys Arg Arg Ile Thr Arg Tyr Asp
 145 150 155 160
 Ala Gln Leu Ile Leu Glu Asn Asn Ser Gly Ile Pro Lys Leu Thr Leu
 165 170 175
 Arg Arg Arg His Asp Ser Ser Ser Lys Thr Asn Asp Gln Glu Asn Asp
 180 185 190
 Gly Met Asn Ser Ser Lys Ile Ser Ile Lys Leu Ser Lys Asp His Asp
 195 200 205
 Asn Asp Asn Asn Leu Tyr Val Ala Lys Leu Asn Asn Gly Phe Asn Ser
 210 215 220
 Gly Ser Gly Ser Ser Ser Thr Lys Leu Lys Ile Gln Leu Lys Arg Asp
 225 230 235 240
 Glu Glu Asn Arg Gly Ser Tyr Thr Glu Gly Leu His Glu Asn Gly Val
 245 250 255
 Cys Cys Ser Asp Pro Leu Ser Leu Leu Glu Ser Arg Met Glu Val Asp
 260 265 270
 Asp Tyr Ser Gln Tyr Glu Glu Glu Ser Thr Asp Asp Ser Ser Ser Ser
 275 280 285
 Glu Gly Asp Glu Glu Glu Asp Asp Tyr Asp Asp Asp Phe Glu Asp Asp
 290 295 300
 Phe Ile Pro Leu Pro Pro Ala Lys Arg Leu Arg Leu Ile Val Gly Lys
 305 310 315 320
 Asp Ser Ile Asp Ile Asp Ile Ser Ser Arg Arg Arg Glu Asp Gln Ser
 325 330 335

Leu Arg Leu Asn Ala
340

<210> 95
<211> 197
<212> PRT
<213> Homo sapiens

<220>
<221> SITE
<222> (179)
<223> Xaa equals any of the naturally occurring L-amino acids

<220>
<221> SITE
<222> (189)
<223> Xaa equals any of the naturally occurring L-amino acids

<220>
<221> SITE
<222> (190)
<223> Xaa equals any of the naturally occurring L-amino acids

<400> 95
Met Gln Phe Ala Trp Gln Ser Tyr Lys Arg Tyr Ala Met Gly Lys Asn
1 5 10 15

Glu Leu Arg Pro Leu Thr Lys Asp Gly Tyr Glu Gly Asn Met Phe Gly
20 25 30

Gly Leu Ser Gly Ala Thr Val Ile Asp Ser Leu Asp Thr Leu Tyr Leu
35 40 45

Met Glu Leu Lys Glu Glu Phe Gln Glu Ala Lys Ala Trp Val Gly Glu
50 55 60

Ser Phe His Leu Asn Val Ser Gly Glu Ala Ser Leu Phe Glu Val Asn
65 70 75 80

Ile Arg Tyr Ile Gly Gly Leu Leu Ser Ala Phe Tyr Leu Thr Gly Glu
85 90 95

Glu Val Phe Arg Ile Lys Ala Ile Arg Leu Gly Glu Lys Leu Leu Pro
100 105 110

Ala Phe Asn Thr Pro Thr Gly Ile Pro Lys Gly Val Val Ser Phe Lys
115 120 125

Ser Gly Asn Trp Gly Trp Ala Thr Ala Gly Ser Ser Ile Leu Ala
130 135 140

Glu Phe Gly Ser Leu His Leu Glu Phe Leu His Leu Thr Glu Leu Ser
145 150 155 160

Gly Asn Gln Val Phe Ala Glu Lys Val Arg Asn Ile Arg Lys Val Leu
165 170 175

Arg Lys Xaa Glu Lys Pro Phe Gly Leu Tyr Ser Asn Xaa Xaa Met Val
180 185 190

Leu Gln Thr Asp Pro
195

<210> 96

<211> 254

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (162)

<223> Xaa equals any of the naturally occurring L-amino acids

<220>

<221> SITE

<222> (170)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 96

Phe	Gly	Thr	Ser	Tyr	Ile	Gly	Gly	Leu	Leu	Ser	Ala	Phe	Tyr	Leu	Thr
1				5					10					15	

Gly	Glu	Glu	Val	Phe	Arg	Ile	Lys	Ala	Ile	Arg	Leu	Gly	Glu	Lys	Leu
			20					25					30		

Leu	Pro	Ala	Phe	Asn	Thr	Pro	Thr	Gly	Ile	Pro	Lys	Gly	Val	Val	Ser
		35					40					45			

Phe	Lys	Ser	Gly	Asn	Trp	Gly	Trp	Ala	Thr	Ala	Gly	Ser	Ser	Ser	Ile
	50					55					60				

Leu	Ala	Glu	Phe	Gly	Ser	Leu	His	Leu	Glu	Phe	Leu	His	Leu	Thr	Glu
65					70					75					80

Leu	Ser	Gly	Asn	Gln	Val	Phe	Ala	Glu	Lys	Val	Arg	Asn	Ile	Arg	Lys
			85						90					95	

Val	Leu	Arg	Lys	Ile	Glu	Lys	Pro	Phe	Gly	Leu	Tyr	Pro	Asn	Phe	Leu
			100					105					110		

Ser	Pro	Val	Ser	Gly	Asn	Trp	Val	Gln	His	His	Val	Ser	Val	Gly	Gly
		115				120						125			

Leu	Gly	Asp	Ser	Phe	Tyr	Glu	Tyr	Leu	Ile	Lys	Ser	Trp	Leu	Met	Ser
	130					135						140			

Gly	Lys	Thr	Asp	Met	Glu	Ala	Lys	Asn	Met	Tyr	Tyr	Glu	Ala	Leu	Glu
145					150					155					160

Ala	Xaa	Arg	Asp	Leu	Leu	Ala	Glu	Cys	Xaa	Ser	Arg	Gly	Ala	Asp	Leu
				165					170					175	

His	Cys	Arg	Val	Ala	Arg	Gly	Asp	Ser	Gly	Pro	Gln	Asp	Gly	Ala	Pro
			180					185					190		

Gly	Leu	Phe	Leu	Arg	Gly	His	Asp	Arg	Pro	Trp	Pro	Glu	Asp	Ala	Lys
		195					200					205			

Glu	Glu	Lys	Arg	Ala	His	Tyr	Arg	Glu	Leu	Ala	Ala	Gln	Ile	Thr	Lys
		210				215					220				

Thr	Cys	His	Glu	Ser	Tyr	Ala	Arg	Ser	Asp	Thr	Lys	Leu	Gly	Pro	Glu
225					230					235					240

Ala	Ser	Gly	Leu	Thr	Pro	Ala	Glu	Arg	Pro	Trp	Pro	Pro	Ser
				245					250				

<210> 97

<211> 97

<212> PRT

<213> Homo sapiens

<400> 97

Met Thr Phe Gln Phe Asn Phe Thr Ile Glu Asp His Leu Glu Asn Glu
 1 5 10 15

Leu Thr Pro Ile Arg Asp Gly Ala Leu Thr Leu Asp Ser Ser Lys Glu
 20 25 30

Leu Ser Val Ser Glu Ser Gln Lys Gly Glu Glu Arg Asp Arg Lys Cys
 35 40 45

Ser Ala Glu Gln Phe Asp Leu Pro Gln Asp His Leu Trp Glu His Lys
 50 55 60

Ser Met Glu Asn Ala Ala Pro Ser Gln Asp Thr Asp Ser Pro Leu Ser
 65 70 75 80

Ala Ala Ser Ser Ser Arg Asn Leu Gly Ala Thr Trp Glu Asn Ser Pro
 85 90 95

Pro

<210> 98

<211> 288

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (277)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 98

Pro His Arg Val Asp Thr Arg Arg Arg Asp Pro Val Pro Arg Ser Arg
 1 5 10 15

Ala Leu Ser His Gly Thr Gly Arg Val Gly Ala Ala Ala Gly Glu Ser
 20 25 30

Ser Arg Ala Pro Arg Cys Trp Ser Gly Ser Arg Pro Arg Ala Pro Ala
 35 40 45

Asp Pro Pro Arg His Arg Pro Leu Leu Cys Leu Ser Arg Arg Gly Ser
 50 55 60

Pro Pro His His Leu Gly Cys Leu Leu Gly Glu Ser Phe Met Gln Leu
 65 70 75 80

Gln Gln Arg Leu Leu Arg Glu Lys Glu Ala Lys Ile Arg Lys Ala Leu
 85 90 95

Asp Arg Leu Arg Lys Lys Arg His Leu Leu Arg Arg Gln Arg Thr Arg
 100 105 110

Arg Glu Phe Pro Val Ile Ser Val Val Gly Tyr Thr Asn Cys Gly Lys
 115 120 125

Thr Thr Leu Ile Lys Ala Leu Thr Gly Asp Ala Ala Ile Gln Pro Arg
 130 135 140

Asp Gln Leu Phe Ala Thr Leu Asp Val Thr Ala His Ala Gly Thr Leu
 145 150 155 160

Pro Ser Arg Met Thr Val Leu Tyr Val Asp Thr Ile Gly Phe Leu Ser
 165 170 175
 Gln Leu Pro His Gly Leu Ile Glu Ser Phe Ser Ala Thr Leu Glu Asp
 180 185 190
 Val Ala His Ser Asp Leu Ile Leu His Val Arg Asp Val Ser His Pro
 195 200 205
 Glu Ala Glu Leu Gln Lys Cys Ser Val Leu Ser Thr Leu Arg Gly Leu
 210 215 220
 Gln Leu Pro Ala Pro Leu Leu Asp Ser Met Val Glu Val His Asn Lys
 225 230 235 240
 Val Asp Leu Val Pro Gly Tyr Ser Pro Thr Glu Pro Asn Val Val Pro
 245 250 255
 Val Ser Ala Leu Arg Gly His Gly Leu Gln Glu Leu Lys Leu Ser Ser
 260 265 270
 Met Arg Arg Phe Xaa Arg Arg Arg Gly Asp Arg Ser Ser Leu Ser Val
 275 280 285

<210> 99
 <211> 94
 <212> PRT
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (61)
 <223> Xaa equals any of the naturally occurring L-amino acids

<220>
 <221> SITE
 <222> (63)
 <223> Xaa equals any of the naturally occurring L-amino acids

<400> 99
 Pro Pro Pro Ile Gly Leu Ala Leu Ala Cys Lys Ser Arg Arg Trp Pro
 1 5 10 15
 Arg Ala Gln Pro Ser Arg Met Ser Pro Gly Pro Pro Leu Trp Glu Arg
 20 25 30
 Arg Gln Ser Tyr Trp Pro Leu Thr Arg Pro Leu Gly Pro Arg Ala Arg
 35 40 45
 Gln Ala Phe Glu Ser Thr Cys Ser Ser Pro Glu Ser Xaa Pro Xaa Arg
 50 55 60
 Ala Ser His Thr Ala Ala Asp Leu Arg Ala Pro Cys Leu Asn Cys Glu
 65 70 75 80
 Phe Phe Leu Gly Asn Pro Leu Lys Arg Lys Gly Tyr Gln Ser
 85 90

<210> 100
 <211> 38

<212> PRT

<213> Homo sapiens

<400> 100

Met Leu Cys Trp Phe Cys Phe Phe Val Leu Leu Phe Phe Phe Phe Phe
 1 5 10 15

Leu Phe Cys Phe Leu Val Met Trp Leu Lys Cys Asn Ser Phe Phe Leu
 20 25 30

Gly His Ile Ser Ala Asn
 35

<210> 101

<211> 60

<212> PRT

<213> Homo sapiens

<400> 101

Met Lys Thr Gly Gly Lys His Ser Val Ile Arg Tyr Phe Ser Asn Ile
 1 5 10 15

Lys Thr Thr Lys Thr Asn Asp Lys Asn Val Tyr Phe Tyr Thr Pro Ala
 20 25 30

Tyr Arg Val Ser Phe Arg Asp Val Tyr Glu Tyr Leu Asn Leu Leu Ile
 35 40 45

Ser Val Leu Met Lys Ala Glu Leu Asn Arg Glu Ser
 50 55 60

<210> 102

<211> 40

<212> PRT

<213> Homo sapiens

<220>

<221> SITE

<222> (6)

<223> Xaa equals any of the naturally occurring L-amino acids

<220>

<221> SITE

<222> (30)

<223> Xaa equals any of the naturally occurring L-amino acids

<400> 102

Trp Met Ser Glu Tyr Xaa Gln Trp Val Phe Leu Ile Ser Leu Arg Ile
 1 5 10 15

Cys Leu Arg Val His Tyr Gln Gly Ile Ser Gly Thr Arg Xaa His Ser
 20 25 30

Leu His Gln Phe Leu Arg Val Leu
 35 40

<210> 103

<211> 228

<212> PRT

<213> Homo sapiens

<400> 103

Met Gly Ala Val Ile Lys Cys Met Met Tyr Tyr Lys Glu Ala Phe Trp

59

1	5	10	15
Lys Lys Lys Asp Tyr Cys Gly Cys Met Ile Ile Glu Asp Glu Asp Ala	20	25	30
Pro Ile Ser Ile Thr Leu Asp Asp Thr Lys Pro Asp Gly Ser Leu Pro	35	40	45
Ala Ile Met Gly Phe Ile Leu Ala Arg Lys Ala Asp Arg Leu Ala Lys	50	55	60
Leu His Lys Glu Ile Arg Lys Lys Lys Ile Cys Glu Leu Tyr Ala Lys	65	70	75
Val Leu Gly Ser Gln Glu Ala Leu His Pro Val His Tyr Glu Glu Lys	85	90	95
Asn Trp Cys Glu Glu Gln Tyr Ser Gly Gly Cys Tyr Thr Ala Tyr Phe	100	105	110
Pro Pro Gly Ile Met Thr Gln Tyr Gly Arg Val Ile Arg Gln Pro Val	115	120	125
Gly Arg Ile Phe Phe Ala Gly Thr Glu Thr Ala Thr Lys Trp Ser Gly	130	135	140
Tyr Met Glu Gly Ala Val Glu Ala Gly Glu Arg Ala Ala Arg Glu Val	145	150	155
Leu Asn Gly Leu Gly Lys Val Thr Glu Lys Asp Ile Trp Val Gln Glu	165	170	175
Pro Glu Ser Lys Asp Val Pro Ala Val Glu Ile Thr His Thr Phe Trp	180	185	190
Glu Arg Asn Leu Pro Ser Val Ser Gly Leu Leu Lys Ile Ile Gly Phe	195	200	205
Ser Thr Ser Val Thr Ala Leu Gly Phe Val Leu Tyr Lys Tyr Lys Leu	210	215	220
Leu Pro Arg Ser	225		

<210> 104

<211> 82

<212> PRT

<213> Homo sapiens

<400> 104

Phe Phe Val Ile Pro Ser Ser Gly Ser Ile Cys Phe Cys Ser Leu Val	1	5	10	15
Thr Val Leu Met Phe Asn Cys Cys Thr Leu Lys Pro Lys Ser Val Thr	20	25	30	
Met His Thr Val Thr Lys Val Leu Gly Leu Gln Ser Cys Leu Leu Tyr	35	40	45	
Lys Glu Asn Phe Lys Cys Cys Cys Lys Leu Thr Ser Tyr Thr Ile Leu	50	55	60	
Asn Phe Leu Ser Ser Pro Leu Phe Leu Pro Thr Asn Gly Ile Ile Met	65	70	75	80

Leu Ala

<210> 105
 <211> 79
 <212> PRT
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (6)
 <223> Xaa equals any of the naturally occurring L-amino acids

<400> 105
 Met Trp Gly Ser Cys Xaa Leu Glu Tyr Tyr Val Ser Pro Pro Ser Ala
 1 5 10 15

Val Phe Ser Glu His Val Cys Cys Pro Trp Trp Glu Arg Gly His Cys
 20 25 30

Ala Val Val His Arg Cys Leu Ser Phe Thr Val Gly Leu Ser Val Cys
 35 40 45

Leu Ser Phe Leu Ser Ala Ala Gln Met Glu Asn Asn Tyr Leu Leu His
 50 55 60

Trp Arg Glu Arg Lys Ser Leu Arg Ile Pro Lys Gly Thr Leu Ala
 65 70 75

<210> 106
 <211> 36
 <212> PRT
 <213> Homo sapiens

<400> 106
 Met Cys Pro Phe Ser Ser Leu His Leu Ala Ala Gly Ile Val Asp Ile
 1 5 10 15

Thr Gly Ala Leu Ala Ala Val Ser Arg Gly Ser Lys Pro His Pro Lys
 20 25 30

Ser Lys Ala Asp
 35

<210> 107
 <211> 35
 <212> PRT
 <213> Homo sapiens

<400> 107
 Met Ala Leu Gly Thr Trp Lys Arg Val Thr Glu Met Gly Arg His Glu
 1 5 10 15

Leu Lys Glu Ala Ile Tyr Asp His Val Ile Cys Asn Met Lys Lys Ala
 20 25 30

Tyr Leu Glu
 35

<210> 108
 <211> 43
 <212> PRT

<213> Homo sapiens

<400> 108

Ile Val Thr Lys Leu Lys Arg Leu Cys Cys Phe Ser Val Met Ser Ala
1 5 10 15

Gly Ile Lys Ala Val Ser Ala Pro Cys Gly Ala Ser Cys Gly Ile Cys
20 25 30

Ser Ser Pro Tyr Pro His Asn Ser Gly Ala Gln
35 40

<210> 109

<211> 74

<212> PRT

<213> Homo sapiens

<400> 109

Arg Pro Thr Arg Pro Pro Cys His Ile Leu Leu Ala Tyr Leu Phe Phe
1 5 10 15

Leu Trp Leu Cys Met Ala Phe Leu Gln Val Leu Ser Arg Tyr His Ser
20 25 30

Ala Asn His Cys Tyr Arg Met Val Thr Ser Phe Val Leu Thr Val Gln
35 40 45

Gln Gln Ile Trp Val Arg Leu Asn Leu Ser Val Asn Phe Phe Phe Trp
50 55 60

Cys Phe Phe Gly Leu Met Thr Val Ser Leu
65 70

<210> 110

<211> 7

<212> PRT

<213> Homo sapiens

<400> 110

Trp Cys Phe Phe Gly Leu Met
1 5

<210> 111

<211> 7

<212> PRT

<213> Homo sapiens

<400> 111

Phe Phe Gly Leu Met Thr Val
1 5

<210> 112

<211> 9

<212> PRT

<213> Homo sapiens

<400> 112

Trp Cys Phe Phe Gly Leu Met Thr Val
1 5

<210> 113

<211> 69
 <212> PRT
 <213> Homo sapiens

<400> 113
 Phe Leu Gln Val Leu Ser Arg Tyr His Ser Ala Asn His Cys Tyr Arg
 1 5 10 15
 Met Val Thr Ser Phe Val Leu Thr Val Gln Gln Gln Ile Trp Val Arg
 20 25 30
 Leu Asn Leu Ser Val Asn Phe Phe Phe Trp Cys Phe Phe Gly Leu Met
 35 40 45
 Thr Val Ser Leu Leu Tyr Pro Cys Phe Ala Cys Asn Asp Ser Cys Met
 50 55 60
 Val Phe Leu Thr Ser
 65

<210> 114
 <211> 167
 <212> PRT
 <213> Homo sapiens

<400> 114
 Ser Trp Gln Thr Glu Glu Lys Thr Cys Asp Leu Val Gly Glu Lys Gly
 1 5 10 15
 Lys Glu Ser Glu Lys Glu Leu Ala Leu Val Lys Arg Leu Lys Pro Leu
 20 25 30
 Phe Asn Lys Ser Phe Glu Ser Thr Val Gly Gln Gly Ser Asp Thr Tyr
 35 40 45
 Ile Tyr Ile Phe Arg Val Cys Arg Glu Ala Gly Asn His Thr Ser Gly
 50 55 60
 Ala Gly Leu Val Gln Ile Asn Lys Ser Asn Gly Lys Glu Thr Val Val
 65 70 75 80
 Gly Arg Leu Asn Glu Thr His Ile Phe Asn Gly Ser Asn Trp Ile Met
 85 90 95
 Leu Ile Tyr Lys Gly Gly Asp Glu Tyr Asp Asn His Cys Gly Lys Glu
 100 105 110
 Gln Arg Arg Ala Val Val Met Ile Ser Cys Asn Arg His Thr Leu Ala
 115 120 125
 Asp Asn Phe Asn Pro Val Ser Glu Glu Arg Gly Lys Val Gln Asp Cys
 130 135 140
 Phe Tyr Leu Phe Glu Met Asp Ser Ser Leu Ala Cys Ser Pro Glu Ile
 145 150 155 160
 Ser His Leu Ser Val Gly Ser
 165

<210> 115
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 115

Ser Trp Gln Thr Glu Glu Lys Thr Cys Asp Leu Val Gly Glu Lys Gly
1 5 10 15Lys Glu Ser Glu Lys Glu Leu Ala Leu Val Lys Arg Leu Lys Pro Leu
20 25 30

<210> 116

<211> 32

<212> PRT

<213> Homo sapiens

<400> 116

Phe Asn Lys Ser Phe Glu Ser Thr Val Gly Gln Gly Ser Asp Thr Tyr
1 5 10 15Ile Tyr Ile Phe Arg Val Cys Arg Glu Ala Gly Asn His Thr Ser Gly
20 25 30

<210> 117

<211> 32

<212> PRT

<213> Homo sapiens

<400> 117

Ala Gly Leu Val Gln Ile Asn Lys Ser Asn Gly Lys Glu Thr Val Val
1 5 10 15Gly Arg Leu Asn Glu Thr His Ile Phe Asn Gly Ser Asn Trp Ile Met
20 25 30

<210> 118

<211> 32

<212> PRT

<213> Homo sapiens

<400> 118

Leu Ile Tyr Lys Gly Gly Asp Glu Tyr Asp Asn His Cys Gly Lys Glu
1 5 10 15Gln Arg Arg Ala Val Val Met Ile Ser Cys Asn Arg His Thr Leu Ala
20 25 30

<210> 119

<211> 39

<212> PRT

<213> Homo sapiens

<400> 119

Asp Asn Phe Asn Pro Val Ser Glu Glu Arg Gly Lys Val Gln Asp Cys

1 5 10 15
 Phe Tyr Leu Phe Glu Met Asp Ser Ser Leu Ala Cys Ser Pro Glu Ile
 20 25 30
 Ser His Leu Ser Val Gly Ser
 35

<210> 120
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 120
 Tyr Ser Cys His Val Gln His
 1 5

<210> 121
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 121
 Pro Tyr Ser Cys His Val Gln His Ser
 1 5

<210> 122
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 122
 Ala Pro Tyr Ser Cys His Val Gln His Ser Ser
 1 5 10

<210> 123
 <211> 58
 <212> PRT
 <213> Homo sapiens

<400> 123
 Ser Cys Asp Phe Trp Ala Val Lys Asn Val Thr Gly Arg Leu Met Val
 1 5 10 15

Gly Leu Arg Trp Trp Asn His Ile Asp Glu Asp Gly Lys Ser His Trp
 20 25 30

Val Phe Glu Ser Arg Lys Glu Ser Ser Gln Glu Asn Lys Thr Val Ser
 35 40 45

Glu Ala Glu Ser Arg Ile Phe Trp Leu Gly
 50 55

<210> 124
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 124
 Gly Glu Ser Gly Val Gly Lys Thr
 1 5

<210> 125
<211> 9
<212> PRT
<213> Homo sapiens

<400> 125
Ile Gly Glu Ser Gly Val Gly Lys Thr
1 5

<210> 126
<211> 9
<212> PRT
<213> Homo sapiens

<400> 126
Gly Glu Ser Gly Val Gly Lys Thr Asn
1 5

<210> 127
<211> 15
<212> PRT
<213> Homo sapiens

<400> 127
Val Val Leu Ile Gly Glu Ser Gly Val Gly Lys Thr Asn Leu Leu
1 5 10 15

<210> 128
<211> 46
<212> PRT
<213> Homo sapiens

<220>
<221> SITE
<222> (34)
<223> Xaa equals any of the naturally occurring L-amino acids

<400> 128
Tyr Tyr Arg Gly Ala Val Gly Ala Leu Leu Val Phe Asp Leu Thr Lys
1 5 10 15

His Gln Thr Tyr Ala Val Val Glu Arg Trp Leu Lys Glu Leu Tyr Asp
20 25 30

His Xaa Glu Ala Thr Ile Val Val Met Leu Val Gly Asn Lys
35 40 45

<210> 129
<211> 17
<212> PRT
<213> Homo sapiens

<400> 129
Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr Arg Gln Arg
1 5 10 15

Tyr

<210> 130
 <211> 288
 <212> PRT
 <213> Homo sapiens

<400> 130

Thr	Arg	Pro	Arg	Val	His	Leu	Ala	Thr	Val	Ser	Ala	Ser	Ala	Ala	Trp
1				5					10					15	
Asp	Ala	Leu	Gly	Leu	Pro	Val	Arg	Ser	His	Met	Gln	Gly	Ser	Thr	Arg
		20						25					30		
Arg	Met	Gly	Val	Met	Thr	Asp	Val	His	Arg	Arg	Phe	Leu	Gln	Leu	Leu
		35					40					45			
Met	Thr	His	Gly	Val	Leu	Glu	Glu	Trp	Asp	Val	Lys	Arg	Leu	Gln	Thr
	50					55					60				
His	Cys	Tyr	Lys	Val	Asp	Arg	Asn	Ala	Thr	Val	Asp	Lys	Leu	Glu	Asp
65					70					75					80
Phe	Ile	Asn	Asn	Ile	Asn	Ser	Val	Leu	Glu	Ser	Leu	Tyr	Ile	Glu	Ile
				85					90					95	
Lys	Arg	Gly	Val	Thr	Glu	Asp	Asp	Gly	Arg	Pro	Ile	Tyr	Ala	Leu	Val
			100					105					110		
Asn	Leu	Ala	Thr	Thr	Ser	Ile	Ser	Lys	Met	Ala	Thr	Asp	Phe	Ala	Glu
		115					120					125			
Asn	Glu	Leu	Asp	Leu	Phe	Arg	Lys	Ala	Leu	Leu	Ile	Ile	Asp	Ser	Glu
	130					135					140				
Thr	Gly	Phe	Ala	Ser	Ser	Thr	Asn	Ile	Leu	Asn	Leu	Val	Asp	Gln	Leu
145					150					155					160
Lys	Gly	Lys	Lys	Met	Arg	Lys	Lys	Glu	Ala	Glu	Gln	Val	Leu	Gln	Lys
				165				170						175	
Phe	Val	Gln	Asn	Lys	Trp	Leu	Ile	Glu	Lys	Glu	Gly	Glu	Phe	Thr	Leu
			180					185					190		
His	Gly	Arg	Ala	Ile	Leu	Glu	Met	Glu	Gln	Tyr	Ile	Arg	Glu	Thr	Pro
		195					200					205			
Asp	Ala	Val	Lys	Ile	Cys	Asn	Ile	Cys	His	Ser	Leu	Leu	Ile	Gln	Gly
	210					215					220				
Gln	Ser	Cys	Glu	Thr	Cys	Gly	Ile	Arg	Met	His	Leu	Pro	Cys	Val	Ala
225					230					235					240
Lys	Tyr	Phe	Gln	Ser	Asn	Ala	Glu	Pro	Arg	Cys	Pro	His	Cys	Asn	Asp
				245					250					255	
Tyr	Trp	Pro	His	Glu	Ile	Pro	Lys	Val	Phe	Asp	Pro	Glu	Lys	Glu	Arg
			260					265					270		
Glu	Ser	Gly	Val	Lys	Ser	Asn	Lys	Lys	Ser	Leu	Arg	Ser	Arg	Gln	His
		275					280					285			

<210> 131
 <211> 7

<212> PRT
 <213> Homo sapiens

<400> 131
 Cys Asn Ile Cys His Ser Leu
 1 5

<210> 132
 <211> 7
 <212> PRT
 <213> Homo sapiens

<400> 132
 Ile Cys Asn Ile Cys His Ser
 1 5

<210> 133
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 133
 Ile Cys Asn Ile Cys His Ser Leu
 1 5

<210> 134
 <211> 277
 <212> PRT
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (147)
 <223> Xaa equals any of the naturally occurring L-amino acids

<400> 134
 Met Gln Gly Ser Thr Arg Arg Met Gly Val Met Thr Asp Val His Arg
 1 5 10 15

Arg Phe Leu Gln Leu Leu Met Thr His Gly Val Leu Glu Glu Trp Asp
 20 25 30

Val Lys Arg Leu Gln Thr His Cys Tyr Lys Val His Asp Arg Asn Ala
 35 40 45

Thr Val Asp Lys Leu Glu Asp Phe Ile Asn Asn Ile Asn Ser Val Leu
 50 55 60

Glu Ser Leu Tyr Ile Glu Ile Lys Arg Gly Val Thr Glu Asp Asp Gly
 65 70 75 80

Arg Pro Ile Tyr Ala Leu Val Asn Leu Ala Thr Thr Ser Ile Ser Lys
 85 90 95

Met Ala Thr Asp Phe Ala Glu Asn Glu Leu Asp Leu Phe Arg Lys Ala
 100 105 110

Leu Glu Leu Ile Ile Asp Ser Glu Thr Gly Phe Ala Ser Ser Thr Asn
 115 120 125

Ile Leu Asn Leu Val Asp Gln Leu Lys Gly Lys Lys Met Arg Lys Lys
 130 135 140

Glu Ala Xaa Gln Val Leu Gln Lys Phe Val Gln Asn Lys Trp Leu Ile
 145 150 155 160
 Glu Lys Glu Gly Glu Phe Thr Leu His Gly Arg Ala Ile Leu Glu Met
 165 170 175
 Glu Gln Tyr Ile Arg Glu Thr Tyr Pro Asp Ala Val Lys Ile Cys Asn
 180 185 190
 Ile Cys His Ser Leu Leu Ile Gln Gly Gln Ser Cys Glu Thr Cys Gly
 195 200 205
 Ile Arg Met His Leu Pro Cys Val Ala Lys Tyr Phe Gln Ser Asn Ala
 210 215 220
 Glu Pro Arg Cys Pro His Cys Asn Asp Tyr Trp Pro His Glu Ile Pro
 225 230 235 240
 Lys Val Phe Asp Pro Glu Lys Glu Arg Glu Ser Gly Val Leu Lys Ser
 245 250 255
 Asn Lys Lys Ser Cys Gly Pro Gly Ser Ile Ser His Arg Ala Leu Leu
 260 265 270
 Arg Gly Trp Leu Pro
 275

<210> 135
 <211> 153
 <212> PRT
 <213> Homo sapiens

<400> 135
 Ile Asn Lys Gln Met Asn Tyr Leu Phe Phe Phe Leu Thr Thr Ser Gly
 1 5 10 15
 Leu Tyr Cys Leu Ser Gly Ser His Gly Ser Asn Val Lys Tyr Ile Val
 20 25 30
 Leu Thr Tyr Phe Asn Cys Ser Trp Ser Leu Thr Ser Pro Gly Phe Arg
 35 40 45
 Asp Val Leu Lys Gly Ser Gln Leu Trp Gln Val Thr Asp Ser Trp Glu
 50 55 60
 Met Glu Arg Thr Lys Glu Tyr Ser Ser Cys Leu Thr Phe Leu Pro Thr
 65 70 75 80
 Ala Asp Ile Val Gln Ala Arg Val Met Glu Glu Leu Asn Leu Leu Ala
 85 90 95
 Ser Gln Ala Ala Pro Ile Pro Thr Ser Gln Cys Thr Ala Pro Pro His
 100 105 110
 Leu Phe Ser Pro Leu Ser Leu Thr Ser Pro Phe Ile Met Ser His Lys
 115 120 125
 Ser Gly Thr Val Gly Ser His Tyr Asn Leu Leu Cys His Arg Asp Ser
 130 135 140
 Ile Phe Leu Ile Ser Asn His Val Ser
 145 150

<210> 136

<211> 118
 <212> PRT
 <213> Homo sapiens

<220>
 <221> SITE
 <222> (12)
 <223> Xaa equals any of the naturally occurring L-amino acids

<400> 136
 Phe Asn Cys Ser Trp Ser Leu Thr Ser Pro Gly Xaa Arg Asp Val Leu
 1 5 10 15
 Lys Gly Ser Gln Leu Trp Gln Val Thr Asp Ser Trp Glu Met Glu Arg
 20 25 30
 Thr Lys Glu Tyr Ser Ser Cys Leu Thr Phe Leu Pro Thr Ala Asp Ile
 35 40 45
 Val Gln Ala Arg Val Met Glu Glu Leu Asn Leu Leu Ala Ser Gln Ala
 50 55 60
 Ala Pro Ile Pro Thr Ser Gln Cys Thr Ala Pro Pro His Leu Phe Ser
 65 70 75 80
 Pro Leu Ser Leu Thr Ser Pro Phe Ile Met Ser His Lys Ser Gly Thr
 85 90 95
 Val Gly Ser His Tyr Asn Leu Leu Cys His Arg Asp Ser Ile Phe Leu
 100 105 110
 Ile Ser Asn His Val Ser
 115

<210> 137
 <211> 337
 <212> PRT
 <213> Homo sapiens

<400> 137
 Arg Thr Asn Leu Lys Glu Ala Ser Asp Ile Lys Leu Glu Pro Asn Thr
 1 5 10 15
 Leu Asn Gly Tyr Lys Ser Ser Val Thr Glu Pro Cys Pro Asp Ser Gly
 20 25 30
 Glu Gln Leu Gln Pro Ala Pro Val Leu Gln Glu Glu Glu Leu Ala His
 35 40 45
 Glu Thr Ala Gln Lys Gly Glu Ala Lys Cys His Lys Ser Asp Thr Gly
 50 55 60
 Met Ser Lys Lys Lys Arg Gln Gly Lys Leu Val Lys Gln Phe Ala Lys
 65 70 75 80
 Ile Glu Glu Ser Thr Pro Val His Asp Ser Pro Gly Lys Asp Asp Ala
 85 90 95
 Val Pro Asp Leu Met Gly Pro His Ser Asp Gln Gly Glu His Ser Gly
 100 105 110
 Thr Val Gly Val Pro Val Ser Tyr Thr Asp Cys Ala Pro Ser Pro Val
 115 120 125
 Gly Cys Ser Val Val Thr Ser Asp Ser Phe Thr Lys Asp Ser Phe Arg

130	135	140
Thr Ala Lys Ser Lys Lys Lys Arg Arg Ile Thr Arg Tyr Asp Ala Gln 145 150 155 160		
Leu Ile Leu Glu Asn Asn Ser Gly Ile Pro Lys Leu Thr Leu Arg Arg 165 170 175		
Arg His Asp Ser Ser Ser Lys Thr Asn Asp Gln Glu Asn Asp Gly Met 180 185 190		
Asn Ser Ser Lys Ile Ser Ile Lys Leu Ser Lys Asp His Asp Asn Asn 195 200 205		
Asn Leu Tyr Val Ala Lys Leu Asn Asn Gly Phe Asn Ser Gly Ser Gly 210 215 220		
Ser Ser Ser Thr Lys Leu Lys Ile Gln Leu Lys Arg Asp Glu Glu Asn 225 230 235 240		
Arg Gly Ser Tyr Thr Glu Gly Leu His Glu Asn Gly Val Cys Cys Ser 245 250 255		
Asp Pro Leu Ser Leu Leu Glu Ser Arg Met Glu Val Asp Asp Tyr Ser 260 265 270		
Gln Tyr Glu Glu Ser Thr Asp Asp Ser Ser Ser Ser Glu Gly Asp Glu 275 280 285		
Glu Glu Asp Asp Tyr Asp Asp Asp Phe Glu Asp Asp Phe Ile Pro Leu 290 295 300		
Pro Pro Ala Lys Arg Leu Arg Leu Ile Val Gly Lys Asp Ser Ile Asp 305 310 315 320		
Ile Asp Ile Ser Ser Arg Arg Arg Glu Asp Gln Ser Leu Arg Leu Asn 325 330 335		
Ala		

<210> 138
 <211> 26
 <212> PRT
 <213> Homo sapiens

<400> 138
 Glu Ala Ala Val Ser Lys Pro Ala Gly Asn Trp Asp Val Ala Gly Asp
 1 5 10 15
 Glu Arg Thr Asp Pro Ser Val Leu Pro Ala
 20 25

<210> 139
 <211> 478
 <212> PRT
 <213> Homo sapiens

<400> 139
 Ala Phe Ala Lys Ser Tyr Leu Gly Asp Thr Ile Glu Gly Thr Pro Ala
 1 5 10 15
 Gly Thr Gly Pro Glu Phe Pro Gly Arg Pro Thr Arg Pro Met Gln Phe
 20 25 30

Ala Trp Gln Ser Tyr Lys Arg Tyr Ala Met Gly Lys Asn Glu Leu Arg
 35 40 45
 Pro Leu Thr Lys Asp Gly Tyr Glu Gly Asn Met Phe Gly Gly Leu Ser
 50 55 60
 Gly Ala Thr Val Ile Asp Ser Leu Asp Thr Leu Tyr Leu Met Glu Leu
 65 70 75 80
 Lys Glu Glu Phe Gln Glu Ala Lys Ala Trp Val Gly Glu Ser Phe His
 85 90 95
 Leu Asn Val Ser Gly Glu Ala Ser Leu Phe Glu Val Asn Ile Arg Tyr
 100 105 110
 Ile Gly Gly Leu Leu Ser Ala Phe Tyr Leu Thr Gly Glu Glu Val Phe
 115 120 125
 Arg Ile Lys Ala Ile Arg Leu Gly Glu Lys Leu Leu Pro Ala Phe Asn
 130 135 140
 Thr Pro Thr Gly Ile Pro Lys Gly Val Val Ser Phe Lys Ser Gly Asn
 145 150 155 160
 Trp Gly Trp Ala Thr Ala Gly Ser Ser Ser Ile Leu Ala Glu Phe Gly
 165 170 175
 Ser Leu His Leu Glu Phe Leu His Leu Thr Glu Leu Ser Gly Asn Gln
 180 185 190
 Val Phe Ala Glu Lys Val Arg Asn Ile Arg Lys Val Leu Arg Lys Ile
 195 200 205
 Glu Lys Pro Phe Gly Leu Tyr Pro Asn Phe Leu Ser Pro Val Ser Gly
 210 215 220
 Asn Trp Val Gln His His Val Ser Val Gly Gly Leu Gly Asp Ser Phe
 225 230 235 240
 Tyr Glu Tyr Leu Ile Lys Ser Trp Leu Met Ser Gly Lys Thr Asp Met
 245 250 255
 Glu Ala Lys Asn Met Tyr Tyr Glu Ala Leu Glu Ala Ile Glu Thr Tyr
 260 265 270
 Leu Leu Asn Val Ser Pro Gly Gly Leu Thr Tyr Ile Ala Glu Trp Arg
 275 280 285
 Gly Gly Ile Leu Asp His Lys Met Gly His Leu Ala Cys Phe Ser Gly
 290 295 300
 Gly Met Ile Ala Leu Gly Ala Glu Asp Ala Lys Glu Glu Lys Arg Ala
 305 310 315 320
 His Tyr Arg Glu Leu Ala Ala Gln Ile Thr Lys Thr Cys His Glu Ser
 325 330 335
 Tyr Ala Arg Ser Asp Thr Lys Leu Gly Pro Glu Ala Phe Trp Phe Asn
 340 345 350
 Ser Gly Arg Glu Ala Val Ala Thr Gln Leu Ser Glu Ser Tyr Tyr Ile
 355 360 365
 Leu Arg Pro Glu Val Val Glu Ser Tyr Met Tyr Leu Trp Arg Gln Thr
 370 375 380

His Asn Pro Ile Tyr Arg Glu Trp Gly Trp Glu Val Val Leu Ala Leu
385 390 395 400

Glu Lys Tyr Cys Arg Thr Glu Ala Gly Phe Ser Gly Ile Gln Asp Val
405 410 415

Tyr Ser Ser Thr Pro Asn His Asp Asn Lys Gln Gln Ser Phe Phe Leu
420 425 430

Ala Glu Thr Leu Lys Tyr Leu Tyr Leu Leu Phe Ser Glu Asp Asp Leu
435 440 445

Leu Ser Leu Glu Asp Trp Val Phe Asn Thr Glu Ala His Pro Leu Pro
450 455 460

Val Asn His Ser Asp Ser Ser Gly Arg Ala Trp Gly Arg His
465 470 475

<210> 140

<211> 4

<212> PRT

<213> Homo sapiens

<400> 140

Asn Val Ser Gly

1

<210> 141

<211> 4

<212> PRT

<213> Homo sapiens

<400> 141

Asn His Ser Asp

1

<210> 142

<211> 8

<212> PRT

<213> Homo sapiens

<400> 142

Gly Tyr Thr Asn Cys Gly Lys Thr

1

5

<210> 143

<211> 10

<212> PRT

<213> Homo sapiens

<400> 143

Val Gly Tyr Thr Asn Cys Gly Lys Thr Thr

1

5

10

<210> 144

<211> 12

<212> PRT

<213> Homo sapiens

<400> 144

Val Val Gly Tyr Thr Asn Cys Gly Lys Thr Thr Leu
 1 5 10

<210> 145

<211> 273

<212> PRT

<213> Homo sapiens

<400> 145

Arg His His Asp Arg Ser Pro Leu Ser Asp Pro Leu Leu Pro Glu Thr
 1 5 10 15

Leu Leu Ala Pro Pro Asp Pro Pro Gly Leu Trp Pro Ala Ala Pro Leu
 20 25 30

Ser Leu Arg Arg Arg Gly Ser Ala Val Thr His Gln Arg Ala Ser Gly
 35 40 45

Arg Gly Trp Gly Gly Gly Ala Gly Met Ser Leu Pro Leu Arg Ala Pro
 50 55 60

Ala Pro Arg Leu Glu Arg Arg Pro Ala Gly Pro Pro Ala Asp Val Phe
 65 70 75 80

Leu Val Pro Lys Arg Val Val Arg Ala Ser Arg Pro Leu Arg Asp Leu
 85 90 95

Arg Ala Ser His Arg Ala Pro Arg Thr Gln Arg Ala Trp Ser Ser Pro
 100 105 110

Leu Thr Pro Ser Pro Ala Gly Thr His Ala Gly Ser Thr His Ser Ala
 115 120 125

Pro Pro Pro Asn Phe Trp Glu Arg Thr Pro Gly Ser Ala Gln Pro Leu
 130 135 140

Ala Phe Gln Lys Pro Leu Tyr Ala Tyr Leu Ile Phe Val Ile Gly Asp
 145 150 155 160

Glu Pro Ser Leu Leu Ser Pro Phe Pro His Thr His Gln Ser Pro Leu
 165 170 175

Ala Ile Pro Ser Pro Ser Ala Ser Pro Pro Pro Ser Cys Ala Pro Ala
 180 185 190

Pro His Ser Pro Pro Pro Ile Gly Leu Ala Leu Ala Cys Lys Ser Arg
 195 200 205

Arg Trp Pro Arg Ala Gln Pro Ser Arg Met Ser Pro Gly Pro Pro Leu
 210 215 220

Trp Glu Arg Arg Gln Ser Tyr Trp Pro Leu Thr Arg Pro Leu Gly Pro
 225 230 235 240

Arg Ala Arg Gln Ala Phe Glu Ser Thr Cys Ser Ser Pro Glu Ser Arg
 245 250 255

Pro Arg Pro Cys Leu Pro His Arg Arg Pro Gln Ser Thr Leu Pro Gln
 260 265 270

Leu

<210> 146

<211> 52
 <212> PRT
 <213> Homo sapiens

<400> 146

Ala Leu Trp Ala Gly Ala Gly Gly Phe Glu Gly Leu Ser Ser Thr Arg
 1 5 10 15
 Ala Gln Arg Ser Cys Gln Trp Pro Val Ala Leu Pro Pro Phe Pro Glu
 20 25 30
 Arg Gly Ser Arg Gly His Pro Gly Arg Leu Gly Pro Gly Pro Pro Ser
 35 40 45
 Ala Leu Ala Ser
 50

<210> 147
 <211> 51
 <212> PRT
 <213> Homo sapiens

<400> 147

Lys Gly Ile Met Leu Cys Trp Phe Cys Phe Phe Val Leu Leu Phe Phe
 1 5 10 15
 Phe Phe Phe Leu Phe Cys Phe Leu Val Met Trp Leu Lys Cys Asn Ser
 20 25 30
 Phe Phe Phe Gly Thr Tyr Phe Cys Gln Leu Lys Thr Arg Arg Ala Gln
 35 40 45
 Leu Phe Phe
 50

<210> 148
 <211> 59
 <212> PRT
 <213> Homo sapiens

<400> 148

Met Lys Thr Gly Gly Lys His Ser Val Ile Arg Tyr Phe Ser Asn Ile
 1 5 10 15
 Lys Thr Thr Lys Thr Asn Asp Lys Asn Val Tyr Phe Tyr Thr Pro Ala
 20 25 30
 Tyr Arg Val Ser Phe Arg Val Tyr Glu Tyr Leu Asn Leu Leu Ile Ser
 35 40 45
 Val Leu Met Lys Ala Glu Leu Asn Arg Glu Ser
 50 55

<210> 149
 <211> 60
 <212> PRT
 <213> Homo sapiens

<400> 149

Pro Gly Lys Pro Lys Ser Ala His Phe Pro Pro Cys Cys Met Phe Ser
 1 5 10 15
 Val Leu Cys Leu Cys Val Cys Ala Arg Gln Arg Asp Arg Leu Phe Val

75

20 25 30

Lys Ser Ala Ser Cys Leu Gly Ile Phe Val Ser His Leu Ala Val Ser
35 40 45

Ser Arg Thr Ile Gln Leu Ala Phe Gln Ala Trp Arg
50 55 60

<210> 150
<211> 39
<212> PRT
<213> Homo sapiens

<220>
<221> SITE
<222> (6)
<223> Xaa equals any of the naturally occurring L-amino acids

<220>
<221> SITE
<222> (29)
<223> Xaa equals any of the naturally occurring L-amino acids

<400> 150
Trp Met Ser Glu Tyr Xaa Gln Trp Val Phe Leu Ile Ser Leu Arg Ile
1 5 10 15

Cys Leu Arg Val His Tyr Gln Gly Ser Gly Thr Arg Xaa His Ser Leu
20 25 30

His Gln Phe Leu Arg Val Leu
35

<210> 151
<211> 37
<212> PRT
<213> Homo sapiens

<400> 151
Arg Lys Lys Lys Ile Cys Glu Leu Tyr Ala Lys Val Leu Gly Ser Gln
1 5 10 15

Glu Ala Leu His Pro His Tyr Glu Glu Lys Asn Trp Cys Glu Glu Gln
20 25 30

Tyr Ser Gly Gly Cys
35

<210> 152
<211> 33
<212> PRT
<213> Homo sapiens

<400> 152
Cys Glu Leu Tyr Ala Lys Val Leu Gly Ser Gln Glu Ala Leu His Pro
1 5 10 15

His Tyr Glu Glu Lys Asn Trp Cys Glu Glu Gln Tyr Ser Gly Gly Cys
20 25 30

Tyr

<210> 153
 <211> 25
 <212> PRT
 <213> Homo sapiens

<400> 153
 Cys Glu Leu Tyr Ala Lys Val Leu Gly Ser Gln Glu Ala Leu His Pro
 1 5 10 15
 Val His Tyr Glu Glu Lys Asn Trp Cys
 20 25

<210> 154
 <211> 109
 <212> PRT
 <213> Homo sapiens

<400> 154
 Gln Leu Leu Leu Leu Pro Pro Lys Ala Pro Arg Asn Pro Phe Leu Pro
 1 5 10 15
 Cys Pro Gly Ser Arg Thr Pro Gly Tyr Ile Trp Lys Val Glu Met Trp
 20 25 30
 Gly Ser Cys Val Leu Glu Tyr Tyr Val Ser Pro Pro Ser Ala Val Phe
 35 40 45
 Ser Glu His Val Cys Cys Pro Trp Trp Glu Arg Gly His Cys Ala Val
 50 55 60
 Val His Arg Cys Leu Ser Phe Thr Val Gly Leu Ser Val Cys Leu Ser
 65 70 75 80
 Phe Leu Ser Ala Ala Gln Met Glu Asn Asn Tyr Leu Leu His Trp Arg
 85 90 95
 Glu Arg Lys Ser Leu Arg Ile Pro Lys Gly Thr Leu Ala
 100 105

<210> 155
 <211> 44
 <212> PRT
 <213> Homo sapiens

<400> 155
 Asp Glu Val Ser Ser Lys Glu Gly Ser Met Cys Pro Ser Ser Leu His
 1 5 10 15
 Leu Ala Ala Gly Ile Val Asp Ile Thr Gly Ala Leu Ala Ala Val Ser
 20 25 30
 Arg Gly Ser Lys Pro His Pro Lys Ser Lys Ala Asp
 35 40

<210> 156
 <211> 186
 <212> PRT
 <213> Homo sapiens

<400> 156
 Gly Gln Arg Gln Ala Leu Cys Pro Gln Leu Ile Leu Glu Ala Ser Arg
 1 5 10 15

Leu	Cys	Glu	Val	Ser	Thr	Ser	Gln	His	Leu	Cys	Ser	Ser	Phe	Glu	Ala
			20					25					30		
Ser	Asn	Cys	Leu	Gly	Lys	Arg	Asp	Arg	Glu	Met	Glu	Ala	Trp	Ile	Arg
		35					40					45			
Ala	Asn	Gln	Pro	Ala	Phe	Leu	Val	Trp	Arg	Ser	Thr	Trp	Pro	Phe	Pro
	50					55					60				
Trp	Ala	Gln	Gly	His	Leu	Lys	His	Cys	Pro	Val	Lys	Leu	Val	Leu	Gly
65					70					75					80
Cys	Pro	Cys	Ala	Trp	Arg	Val	Leu	Lys	Leu	Thr	Phe	Gln	Ile	Pro	Arg
				85					90					95	
Glu	Gln	Gly	Glu	Ile	Ser	Arg	Met	Ser	Ile	Ala	Ala	Lys	Lys	Cys	Leu
			100					105					110		
Gly	Gly	Leu	Pro	Leu	Leu	Thr	Pro	His	Leu	Ala	Ala	Asp	Gln	His	Ser
		115					120					125			
Ile	Leu	Asn	Thr	Leu	Arg	Ala	Pro	Ser	Met	Ala	Phe	Asp	Arg	Thr	Lys
	130					135					140				
Ser	Pro	Gly	Val	Val	Thr	Glu	Asn	Arg	Ser	Cys	Ala	Val	Thr	Ala	Met
145					150					155					160
Phe	Pro	Pro	Gly	Arg	Gln	Lys	Leu	Lys	Ser	Pro	Lys	Arg	Thr	Ser	Phe
				165					170					175	
Ser	Ser	Ala	Ala	Asp	Glu	Trp	His	Arg	Tyr						
			180					185							

```
<210> 157
<211> 102
<212> PRT
<213> Homo sapiens
```

```

<400> 157
Val Ala Ser Ile Leu Lys Ala Ala Pro Asn Arg Gln Ile Leu Pro Leu
  1              5              10              15
Phe Leu Lys His His His Val Gly Glu Pro Ser Glu Gly Trp Ala Thr
      20              25              30
Ser Gln Asp Ser Leu Leu Gly Gly Leu Gly Tyr Leu Gly Val Leu Pro
      35              40              45
His Asn Val Gln Gly Asp Ile Val Thr Lys Leu Lys Arg Leu Cys Cys
      50              55              60
Phe Ser Val Met Ser Ala Gly Ile Lys Ala Val Ser Ala Pro Cys Gly
  65              70              75              80
Ala Ser Cys Gly Ile Cys Ser Ser Pro Tyr Pro His Asn Ser Gly Ala
      85              90              95
Gln Gly Pro Gly Leu Val
      100

```

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US00/19666

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) : C07K 14/47; C12N 5/10, 5/16, 15/12, 15/63, 15/64

US CL : Please See Extra Sheet.

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 530/350; 536/23.1, 23.5, 24.3, 24.31; 435/69.1, 71.1, 71.2, 471, 325, 252.3, 254.11, 320.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
NONE

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X --- A	WO 92/05256 A1 (GENETICS INSTITUTE, INC. THE WISTAR INSTITUTE) 02 April 1992 (02/04/92), see entire document, especially pages 17-21.	1-3, 7-11, 14-16 ----- 4-6, 12, 21

☐ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

* Special categories of cited documents:	*T	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
*A		document defining the general state of the art which is not considered to be of particular relevance
*E	*X	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
*L		document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
*O	*Y	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
*P	*A	document published prior to the international filing date but later than the priority date claimed
		document member of the same patent family

Date of the actual completion of the international search

22 AUGUST 2000

Date of mailing of the international search report

07 SEP 2000

Name and mailing address of the ISA/US
Commissioner of Patents and Trademarks
Box PCT
Washington, D.C. 20231

Facsimile No. (703) 305-3230

Authorized officer

PREMA WERTZ

Telephone No. (703) 308-0196

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US00/19666

Box I Observations where certain claims were found unsearchable (Continuation of Item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of Item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

Please See Extra Sheet.

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. ☒ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
1-12, 14-16, 21

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US00/19666

A. CLASSIFICATION OF SUBJECT MATTER:

US CL :

530/350; 536/23.1, 23.5, 24.3, 24.31; 435/69.1, 71.1, 71.2, 471, 325, 252.3, 254.11, 320.1

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING

This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Group I, claims 1-12, 14-16, 21, drawn to a nucleic acid of SEQ ID NO:11 encoding a protein of SEQ ID NO:60, a vector, a host cell, a method of making the protein and the protein of SEQ ID NO:60.

Group II, claim 13, drawn to an antibody that binds the protein of SEQ ID NO:60.

Group III, claim 17, drawn to a method of treating a condition comprising administering the protein of SEQ ID NO:60.

Group IV, claim 18, drawn to a method of diagnosing a pathological condition using the polynucleotide encoding a protein of SEQ ID NO:60.

Group V, claim 19, drawn to a method of diagnosing a pathological condition by determining the amount of protein of SEQ ID NO:60.

Group VI, claim 20, drawn to a method of identifying a binding partner of the protein of SEQ ID NO:60.

Group VII, claims 22-23, drawn to a method of identifying an activity in a biological assay.

Group VIII, claim 17, drawn to a method for treating a condition comprising administering the nucleic acid of SEQ ID NO:11.

The inventions listed as Groups I-VIII do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons:

Pursuant to 37 C.F.R. § 1.475 (d), the ISA/US considers that where multiple products and processes are claimed, the main invention shall consist of the first invention of the category first mentioned in the claims and the first recited invention of each of the other categories related thereto. Accordingly, the main invention (Group I) comprises the first-recited product, a nucleic acid encoding a protein of SEQ ID NO:60, a vector, a host cell, a method of making the protein of SEQ ID NO:60, and the protein of SEQ ID NO:60. Further pursuant to 37

C.F.R. § 1.475 (d), the ISA/US considers that any feature which the subsequently recited products and methods share with the main invention does not constitute a special technical feature within the meaning of PCT Rule 13.2 and that each of such products and methods accordingly defines a separate invention.

This application contains claims directed to more than one species of the generic invention. These species are deemed to lack Unity of Invention because they are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for more than one species to be searched, the appropriate additional search fees must be paid. The species are as follows:

the polynucleotides set forth in SEQ ID NO:11-59 encoding the polypeptides set forth in SEQ ID NO:60-108.